



SEQUENCE LISTING

<110> Rosenberg, Eugene

Ron, Eliora

Orr, Elisha

Paitan, Yossi

<120> GENE CLUSTER

<130> 2290.00101

<140> 09/710,262

<141> 2000-11-10

<160> 20

<170> PatentIn Ver. 2.1

<210> 1

<211> 2392

<212> DNA

<213> Myxococcus xanthus

<400> 1

Val Asp Pro Ala Arg Leu Thr Arg Ala Trp Glu Gly Leu Leu Glu Arg
1 5 10 15

Tyr Pro Leu Leu Ala Gly Ala Ile Arg Val Glu Gly Thr Glu Pro Val
20 25 30

Ile Val Pro Ser Gly Gln Val Ser Ala Glu Val His Glu Val Pro Ser
35 40 45

Val Ser Asp Ser Ala Leu Val Ala Thr Leu Arg Ala Ser Ala Lys Val
50 55 60

Pro Phe Asp Leu Ala Cys Gly Pro Leu Ala Arg Leu His Leu Tyr Ser
65 70 75 80

Arg Ser Glu His Glu His Val Leu Leu Cys Phe His His Leu Val
85 90 95

Leu Asp Gly Ala Ser Val Ala Pro Leu Leu Asp Ala Leu Arg Glu Arg
100 105 110

Tyr Ala Gly Thr Glu Ala Lys Ala Gly Leu Leu Glu Val Pro Ile Val
115 120 125

Ala Pro Tyr Arg Ala Ala Val Glu Trp Glu Gln Leu Ala Ile Gly Gly
130 135 140

Asp Glu Gly Arg Arg His Leu Asp Tyr Trp Arg His Val Leu Ala Thr
145 150 155 160

RECEIVED

APR 19 2002

TECH CENTER 1600/2900

Tyr Glu Gly His Ser Leu Ser Tyr Gln Ala Leu Asp Thr Arg Ser Arg
450 455 460

Glu Ile Ala Ala His Leu Lys Ser Phe Gly Val Lys Pro Gly Ala Leu
465 470 475 480

Val Gly Ile Tyr Leu Asp Arg Ser Ala Glu Leu Val Ala Ala Met Leu
485 490 495

Gly Val Leu Ser Ala Gly Ala Ala Tyr Val Pro Leu Asp Pro Val His
500 505 510

Pro Glu Asp Arg Leu Arg Tyr Met Leu Glu Asp Ser Gly Val Val Val
515 520 525

Val Leu Ala Arg Gln Ala Ser Arg Asp Lys Val Ala Ala Ile Ala Gly
530 535 540

Ala Ser Cys Lys Val Cys Val Leu Glu Asp Val Lys Ala Gly Ala Thr
545 550 555 560

Ser Ala Pro Ala Gly Thr Ser Pro Asn Gly Leu Ala Tyr Val Ile Tyr
565 570 575

Thr Ser Gly Ser Thr Gly Arg Pro Lys Gly Val Met Ile Pro His Arg
580 585 590

Gly Val Val Asn Phe Leu Leu Cys Met Arg Arg Thr Leu Gly Leu Lys
595 600 605

Arg Thr Asp Ser Leu Leu Ala Val Thr Thr Tyr Cys Phe Asp Ile Ala
610 615 620

Ala Leu Glu Leu Leu Pro Leu Cys Ala Gly Ala Gln Val Ile Ile
625 630 635 640

Ala Ser Ala Glu Thr Val Arg Asp Ala Gln Ala Leu Lys Arg Ala Leu
645 650 655

Arg Thr His Arg Pro Thr Leu Met Gln Ala Thr Pro Ala Thr Trp Thr
660 665 670

Leu Leu Phe Gln Ser Gly Trp Glu Asn Ala Glu Arg Val Arg Leu
675 680 685

Cys Gly Gly Glu Ala Leu Pro Glu Ser Leu Lys Ala His Phe Val Arg
690 695 700

Thr Ala Ser Asp Val Trp Asn Met Phe Gly Pro Thr Glu Thr Thr Ile
705 710 715 720

Trp Ser Thr Met Ala Lys Val Ser Ala Ser Arg Pro Val Thr Ile Gly
725 730 735

Lys Pro Ile Asp Asn Thr Gln Val Tyr Val Leu Asp Asp Arg Met Gln
740 745 750

Pro Val Pro Ile Gly Val Pro Gly Glu Leu Trp Ile Ala Gly Ala Gly
755 760 765

Val Ala Cys Gly Tyr Leu Asn Arg Pro Ala Leu Thr Ala Glu Arg Phe
770 775 780

Val Ser Asn Pro Phe Thr Pro Gly Thr Thr Leu Tyr Arg Thr Gly Asp
785 790 795 800

Leu Ala Arg Trp Arg Ala Asp Gly Glu Val Glu Tyr Leu Gly Arg Leu
805 810 815

Asp His Gln Val Lys Val Arg Gly Phe Arg Ile Glu Met Gly Glu Ile
820 825 830

Glu Ala Gln Leu Ala Gly His Pro Ser Val Lys Asn Cys Ala Val Val
835 840 845

Ala Lys Glu Leu Asn Gly Thr Ser Gln Leu Val Ala Tyr Cys Gln Pro
850 855 860

Ala Gly Thr Ser Phe Asp Glu Glu Ala Ile Arg Ala His Leu Arg Lys
865 870 875 880

Phe Leu Pro Asp Tyr Met Val Pro Ala His Val Phe Ala Val Asp Ala
885 890 895

Ile Pro Leu Ser Gly Asn Gly Lys Val Asp Arg Gly Gln Leu Met Ala
900 905 910

Arg Pro Val Val Thr Arg Arg Lys Thr Ser Ala Val His Ala Arg Ser
915 920 925

Pro Val Glu Ala Thr Leu Val Glu Leu Trp Lys Asn Val Leu Gln Val
930 935 940

Asn Glu Val Gly Val Glu Asp Arg Phe Phe Glu Val Gly Gly Asp Ser
945 950 955 960

Val Leu Ala Ala Val Leu Val Glu Glu Met Asn Arg Arg Phe Asp Thr
965 970 975

Arg Leu Ala Val Thr Asp Leu Phe Lys Tyr Val Asn Ile Arg Asp Met
980 985 990

Ala Arg His Met Glu Gly Ala Thr Ala Gln Ala Arg Thr Gly Ala Thr
995 1000 1005

Glu Pro Ala Arg Glu Asp Thr Ala Ser Glu Arg Asp Tyr Glu Gly Ser
1010 1015 1020

Leu Ala Val Ile Gly Ile Ser Cys Gln Leu Pro Gly Ala Ala Asp Pro
1025 1030 1035 1040

Trp Arg Phe Trp Lys Asn Leu Arg Glu Gly Arg Asp Ser Val Val Ala
1045 1050 1055

Tyr Arg His Glu Glu Leu Arg Glu Leu Gly Val Pro Glu Glu Val Leu
1060 1065 1070

Arg Asp Ser Arg Tyr Val Ala Val Arg Ser Ser Ile Glu Asp Lys Glu
1075 1080 1085

Cys Phe Asp Pro His Phe Phe Gly Leu Thr Ala Arg Asp Ala Ser Phe
1090 1095 1100

Met Asp Pro Gln Phe Arg Leu Leu Leu Met His Ala Trp Lys Ala Val
1105 1110 1115 1120

Glu Asp Ala Ala Thr Thr Pro Glu Arg Leu Gly Pro Cys Gly Val Phe
1125 1130 1135

Met Thr Ala Ser Asn Ser Phe Tyr His Gln Gly Ser Pro Gln Phe Pro
1140 1145 1150

Ala Asp Gly Gln Pro Val Leu Arg Thr Ala Glu Glu Tyr Val Leu Trp
1155 1160 1165

Val Leu Ala Gln Ala Gly Ser Ile Pro Thr Met Val Ser Tyr Lys Leu
1170 1175 1180

Gly Leu Lys Gly Pro Ser Leu Phe Val His Thr Asn Cys Ser Ser Ser
1185 1190 1195 1200

Leu Ser Ala Leu Tyr Val Ala Gln Gln Ala Ile Ala Ala Gly Asp Cys
1205 1210 1215

Gln Thr Ala Leu Val Gly Ala Ala Thr Val Phe Pro Ser Ala Asn Leu
1220 1225 1230

Gly Tyr Leu His Gln Arg Gly Leu Asn Phe Ser Ser Ala Gly Arg Val
1235 1240 1245

Lys Ala Phe Asp Ala Ala Ala Asp Gly Met Ile Ala Gly Glu Gly Val
1250 1255 1260

Ala Val Leu Val Val Lys Asp Ala Ala Ala Val Arg Asp Gly Asp
1265 1270 1275 1280

Pro Ile Tyr Cys Leu Val Arg Lys Val Gly Ile Asn Asn Asp Gly Gln
1285 1290 1295

Asp Lys Val Gly Leu Tyr Ala Pro Ser Ala Thr Gly Gln Ala Glu Val
1300 1305 1310

Ile Arg Arg Leu Phe Asp Arg Thr Gly Ile Asp Pro Ala Ser Ile Gly
1315 1320 1325

Tyr Val Glu Ala His Gly Thr Gly Thr Leu Leu Gly Asp Pro Val Glu
1330 1335 1340

Val Ser Ala Leu Ser Glu Ala Phe Arg Thr Phe Thr Asp Arg Arg Gly
1345 1350 1355 1360

Tyr Cys Arg Leu Gly Ser Val Lys Ser Asn Leu Gly His Leu Asp Thr
1365 1370 1375

Val Ala Gly Leu Ala Gly Leu Ile Lys Thr Ala Leu Ser Leu Arg Gln
1380 1385 1390

Gly Glu Val Pro Pro Thr Leu His Val Thr Gln Val Asn Pro Lys Leu
1395 1400 1405

Glu Leu Thr Asp Ser Pro Phe Val Ile Ala Asp Arg Leu Ala Pro Trp
1410 1415 1420

Pro Ser Leu Pro Gly Pro Arg Arg Ala Ala Val Ser Ala Phe Gly Leu
1425 1430 1435 1440

Gly Gly Thr Asn Thr His Ala Ile Leu Glu His Tyr Pro Arg Asp Ser
1445 1450 1455

Arg Pro Arg Glu Arg Ser Gln Arg Ser Asn Ala Val Arg Ala Val Ala
1460 1465 1470

Pro Phe Ser Ala Arg Thr Leu Glu Ala Leu Lys Asp Asn Leu Arg Ala
1475 1480 1485

Leu Leu Asp Phe Leu Glu Asp Pro Ala Ser Ala Glu Val Ala Leu Ala
1490 1495 1500

Asp Ile Thr Tyr Thr Leu Gln Val Gly Arg Val Ala Met Pro Glu Arg
1505 1510 1515 1520

Met Val Val Thr Ala Ser Thr Arg Asp Glu Leu Val Glu Gly Leu Arg
1525 1530 1535

Arg Gly Ile Ala Thr Val Gly Gly Ala His Val Gly Thr Val Val Asp
1540 1545 1550

Thr Ser Pro Ser Val Asp Ala Asp Ala Arg Ala Val Ala Glu Ala Trp
1555 1560 1565

Ala Thr Gly Asp Ser Ile Asp Trp Asp Ser Leu His Gly Asp Val Lys
1570 1575 1580

Pro Ala Arg Val Ser Leu Pro Thr Tyr Gln Phe Ala Lys Glu Arg Tyr
1585 1590 1595 1600

Gly Leu Ser Pro Ala His Ser Val Ala Asn Ser Ser Lys Thr His Pro
1605 1610 1615

Asp Ala Gly Val Pro Leu Phe Val Pro Thr Trp Gln Pro Trp Ser Glu
1620 1625 1630

Gly Ala Ser Asn Ala Ser Leu Ala Leu Arg His Leu Val Val Leu Cys
1635 1640 1645

Glu Pro Leu Asp Ala Leu Gly Ala Glu Gly Ala Ser Ala Leu Ala Ser
1650 1655 1660

Thr Leu Ala Asp Arg Arg Ile Glu Val Val Arg Thr Ser Ser Pro Ser
1665 1670 1675 1680

Ala Arg Leu Asp Ala Arg Phe Met Ala His Ala Ser Ala Val Phe Glu
1685 1690 1695

Arg Val Lys Ala Leu Leu Ser Glu Arg Leu Thr Ala Pro Val Thr Leu
1700 1705 1710

Gln Val Leu Val Pro Glu Glu Arg Asp Ala Leu Ala Leu Ser Gly Leu
1715 1720 1725

Gly Ser Leu Leu Arg Ser Val Ser Gln Glu Asn Pro Leu Val Arg Gly
1730 1735 1740

Gln Leu Ile Arg Val Gln Gly Ser Val Ser Ala Ser Ala Leu Val Asp
1745 1750 1755 1760

Val Leu Val Lys Ser Ala Arg Ala Gly Asp Val Thr Asp Ser Arg Tyr
1765 1770 1775

His Ala Gly Gln Leu Ser Arg Cys Glu Trp Arg Glu Ala Arg Val Ala
1780 1785 1790

Lys Gly Asp Ala Ser Arg Phe Trp Arg Glu Asp Gly Val Tyr Val Ile
1795 1800 1805

Ser Gly Gly Thr Gly Ala Leu Ala Arg Leu Phe Val Ala Glu Ile Gly
1810 1815 1820

Lys Arg Ala Thr Arg Ala Thr Val Ile Leu Val Ala Arg Ala Ser Ser
1825 1830 1835 1840

Ala Glu Ala Val Asp Gly Gly Asn Gly Leu Arg Val Arg His Leu Pro
1845 1850 1855

Val Asp Val Thr Gln Pro Asn Asp Val Asn Ala Phe Val Ala Thr Val
1860 1865 1870

Leu Arg Glu His Gly Arg Ile Asp Gly Val Ile His Ala Ala Gly Ile
1875 1880 1885

Arg Arg Asp Asn Tyr Leu Leu Asn Lys Pro Val Ala Glu Met Gln Ala
1890 1895 1900

Val Leu Ala Pro Lys Val Val Gly Leu Val Asn Leu Asp His Ala Thr
1905 1910 1915 1920

Arg Glu Leu Pro Leu Asp Phe Phe Val Thr Phe Ser Ser Leu Ala Ala
1925 1930 1935

Phe Gly Asn Ala Gly Gln Ser Asp Tyr Ala Ala Ala Asn Gly Phe Met
1940 1945 1950

Asp Gly Phe Ala Glu Ser Arg Ala Ala Leu Val Asn Ala Gly Gln Arg
1955 1960 1965

Gln Gly Arg Thr Val Ser Ile Arg Trp Pro Leu Trp Glu Asn Gly Gly
1970 1975 1980

Met Gln Leu Asp Ser Arg Ser Arg Glu Val Leu Met Gln Arg Thr Gly
1985 1990 1995 2000

Met Ala Ala Leu Gly Asp Glu Ala Gly Leu Gly Ala Phe Tyr Arg Ala
2005 2010 2015

Leu Glu Leu Gly Ser Pro Gly Val Ala Val Trp Thr Gly Glu Ala Gln
2020 2025 2030

Arg Phe Arg Glu Leu Ser Val Ser Val Ser Pro Ala Pro Pro Pro His
2035 2040 2045

Gln Val Ala Leu Asp Ala Val Val Ser Ile Thr Glu Lys Val Glu Thr
2050 2055 2060

Lys Leu Lys Ala Leu Phe Ser Glu Val Thr Arg Tyr Glu Glu Arg Arg
2065 2070 2075 2080

Ile Asp Ala Arg Gln Pro Met Glu Arg Tyr Gly Ile Asp Ser Ile Ile
2085 2090 2095

Ile Thr Gln Met Asn Gln Ala Leu Glu Gly Pro Tyr Asn Ala Leu Ser
2100 2105 2110

Lys Thr Leu Phe Phe Glu Tyr Arg Thr Leu Ala Glu Val Ser Gly Tyr
2115 2120 2125

Leu Ala Glu His Arg Ala Glu Glu Ser Ala Lys Trp Val Ala Ala Pro
2130 2135 2140

Gly Glu Asn Ser Ser Val Ile Gln Glu Ala Arg Pro Pro Arg Ala
2145 2150 2155 2160

Asp Ala Thr His Arg Ala Pro Arg Ala Asp Glu Pro Ile Ala Val Ile
2165 2170 2175

Gly Met Ser Gly Arg Tyr Pro Gly Ala Glu Asn Leu Thr Glu Phe Trp
 2180 2185 2190
 Glu Arg Leu Ser Arg Gly Asp Asp Cys Ile Thr Glu Ile Pro Pro Glu
 2195 2200 2205
 Arg Trp Ser Leu Asp Gly Phe Phe Tyr Pro Asp Lys Lys His Ala Ala
 2210 2215 2220
 Ala Arg Gly Met Ser Tyr Ser Lys Trp Gly Gly Phe Leu Gly Gly Phe
 2225 2230 2235 2240
 Ala Asp Phe Asp Pro Leu Phe Phe Asn Ile Ser Pro Arg Glu Ala Thr
 2245 2250 2255
 Ser Met Asp Pro Gln Glu Arg Leu Phe Leu Gln Ser Cys Trp Glu Val
 2260 2265 2270
 Leu Glu Asp Ala Gly Tyr Thr Arg Asp Ser Leu Ala Gln Arg Phe Gly
 2275 2280 2285
 Ser Ala Val Gly Val Phe Ala Gly Ile Thr Lys Thr Gly Tyr Glu Leu
 2290 2295 2300
 Tyr Gly Ala Glu Leu Glu Gly Arg Asp Ala Ser Val Arg Pro Tyr Thr
 2305 2310 2315 2320
 Ser Phe Ala Ser Val Ala Asn Arg Val Ser Tyr Leu Leu Asp Leu Lys
 2325 2330 2335
 Gly Pro Ser Met Pro Val Asp Thr Met Cys Ser Ala Ser Leu Thr Ala
 2340 2345 2350
 Val His Met Ala Cys Glu Ala Leu Gln Arg Gly Ala Cys Val Met Ala
 2355 2360 2365
 Ile Ala Gly Gly Val Asn Leu Tyr Val His Pro Ser Ser Tyr Val Ser
 2370 2375 2380
 Leu Ser Gly Gln Gln Met Leu Ser
 2385 2390

<210> 2

<211> 7178

<212> DNA

<213> Myxococcus xanthus

<400> 2

gtcgacccgg cgaggctgac cggggcctgg gaaggactgc tcgaacggta tccgctgctc 60
 gctggcgca ttgcgtcga aggcacggag cgggtcatcg tccccagtgg gcaggtctcc 120
 gccgagggtcc acgagggtcc atcggctcc gattcagcac tggtggcgac cctgcgcgccc 180
 tccgcgaagg tgccattcga tctgcctgt ggaccgctcg ctggctgca cctgtactcg 240
 cggtcggagc acgagcatgt cctgctgtg tgctccacc acctgtgtcg cgtatggggca 300

tccgtggcgc cctgctcg a cgcctccgg gagcgttacg ccgggaccga ggcgaaggcg 360
ggctgtctcg agggtccgt cgtcgtccct taccgcgcgg cctgtggatgt ggagcagctc 420
ccatggag gcatgggg acggccac ctcgactact ggccgcacgt gtggccacg 480
ccgttcctc cgccgttga tttcaacg gaccggcctc gtcggccac ggggctggac 540
tcggagggag caacgcac tgcagagggtg cccaccggc aagcattgcg actgcgcgag 600
ttcgctggg cacagcaatg gagectgcg accgttgc tgggtctta ctacgcctg 660
tttcatggc acacgcgca ggacgcgtg tgggtcgca tccacccat gggcggccc 720
cgggcggaaac tggcgcacggc gattgggtac ttgcgttcaacg tggatggcgt ggcgcgcgg 780
ggcctggggc agcactcgtt cggctcgctg ctgcggccacc tccacggactc ggtcatcgat 840
ggcctggagc atgcccacta tccctcccg ctagtggatgttga aggacactcg gctgtcaat 900
gggcccggagg aggcgcctgg cttccagacg atgttccactt tccagacgc tcaactgacg 960
agcgtccgc caaggccgg a cgcctccgg ggcaggatgtg ggccgggtgc cggagcttga gcccgtcgac 1020
tgcgtccatc aggaaggcgc ctaccgcgtg gagcttgaag tgggtggatgg cgccaaggc 1080
ctcagctgtc atttcaatgtt ctagcgcgg ctgtacgagg cggacacggt cgaacggatg 1140
gcccgtcagt tggcgtcgcc cgcggaccag gtcggggatgt ggggtggatgt tccgttgcg 1200
gcactgtctg ggctcgacga cgaagagcgc cgcacgcctc tccgcgtactg gaatgccacg 1260
gccacgcgt tccctcgagga cctggcggtt cacgagctct tccagcggca gggccgggag 1320
acccccagacg ccatggctgtt gagctacgag gggcactcgc tcagctatca ggcgttggat 1380
acgcggagcc gcgagatgtc ggccgttcaacg aagacgttgc ggcgtcaagcc tggggcgctc 1440
gtgggcatct acctggaccc gtcgcggag ctgggtggcg ctagtgcgtt tggctgtcc 1500
gctggcgccg cctacgtacc cctggaccccg gtgcaccccg aggacccgt ggggtacatg 1560
ctggaggaca gtggcggtt ggtcgctgt gcccgtcagg cctcgccggaa caaggtcgcc 1620
gccatggccg gaggctccgtt caagggtgtc gtgtcgagg acgtcaaggc tggggccacg 1680
tccgcgcgg cgggaaccc accgaacgg a ctcgttgc tcatctacac gtcggggagc 1740
acggggccgc ccaagggcgtt gatgttccc catcgccggg tggcaactt ctcctgtgc 1800
atgcgcagga cgcgtggccctt gaagcgcacg gattcgctgt tggcgttac gacgtactgc 1860
ttcgacatcg cggcgctcg a ctcgttgc tccgttgc tggggggca ggtcatcatc 1920
gctcgccgg agacgggtcg ggatgcgcag gctgttgc gggcgctcg caccatcg 1980
cccacgttga tgcaggcgac gcccgcgacc tggacactgt tggcttgc tggctggag 2040
aacgcggagc ggggttcaatgttgcgtt ggagaagcgc tggccggatgt gtcgaaggcc 2100
cacttcgttc gcacccgcgag ctagtgcgtt aacatgttgc gcccacccga gacgaccatc 2160
tggcgtacga tggcgttgcgtt ctcgttgcgtt ctagtggaaa gccgtatcgac 2220
aacacgcagg tctacgtgtt ggacgaccgg atgcagccgg tggccatcg tggccggcc 2280
gagctgttga ttgcggccg gggcggttacc tcaacccggc ggcgttgcacc 2340
ggcgacgtt tgggttccaa tccgttacgtt cccggccacgtt cgttccatcg gacgggggac 2400
ctggcgccgtt ggcgtgttgcgtt cgggtggatgtt ggttccatcg tggctggatgt 2460
aagggtgcgtt gtcgttgcgtt ctcgttgcgtt ctagtggaaa gccgtatcgac 2520
agcgttgaaga actgttgcgtt ggtggccaa ggttgcgtt gacgttgcgtt gtcgttgcgtt 2580
tactgtcaggc cccggggaaac ggttgcgtt gggggccaa tccgttgcgtt ctcgttgcgtt 2640
ttccctcccg actacatgtt cccggccac ttcgttgcgtt tggatgttgcgtt tccgttgcgtt 2700
ggcaatggca aggttgcgtt gggccagctg atggccaggc cgggtggatgttgcgttccggccggaaag 2760
acatcccgccg tccatggcccg ttcgttgcgtt gggccaccc tccgttgcgtt tggatgttgcgttccggccggaaac 2820
gtgtccagg tcaacgttgcgtt ggggttgcgtt gatgttgcgtt tccgttgcgtt tggatgttgcgttccggccggaaac 2880
gtgtccagg cctgttgcgtt ggggttgcgtt gatgttgcgtt tccgttgcgtt tggatgttgcgttccggccggaaac 2940
accgttgcgtt tcaacgttgcgtt caatatttcgtt gacgttgcgtt gggccggccggaaac 3000
gcccgttgcgtt gtcgttgcgtt ctcgttgcgtt tccgttgcgtt tggatgttgcgttccggccggaaac 3060
tacggggca gtcgttgcgtt ctcgttgcgtt tccgttgcgtt tggatgttgcgttccggccggaaac 3120
tggcgttgcgtt ggaaggaaactt gtcgttgcgtt gggccaggc ggttgcgttccggccggaaac 3180
gaacgttgcgtt gtcgttgcgtt gggccaggc ggttgcgttccggccggaaac 3240
cggttgcgttccggccggaaac 3300
gacgttgcgtt tccgttgcgtt gggccaggc ggttgcgttccggccggaaac 3360
gaagacgcgg cgcgttgcgtt gggccaggc ggttgcgttccggccggaaac 3420
aacacgttgcgtt atcaccaggc ttcgttgcgtt tccgttgcgttccggccggaaac 3480
accggccggaaac aatacgttgcgtt gggccaggc ggttgcgttccggccggaaac 3540

tcstacaagc tcggcttcaa gggccgagc ctgttcgtcc acaccaactg ctcgtcatcc 3600
ctgtccgcgc tgcacgtggc tcagcaggcc atcgcacgccc gagactgcca gacggcgctg 3660
gtggggggcc ccacggctt ccttcggcg aacttgggtt atctgcacca ggggggctc 3720
aacttctcca ggcggggcc ggtcaaggcc ttgcacgccc cggcggacgg catgattgcc 3780
ggtgaagggt tcgcgtgtc ggggtgaag gacgcgcgc cggcgggtcg cgtatggc 3840
ccaatctact gcctcggtc gaaagggtggg atcaacaacg acggccagga caagggtggg 3900
ttatacgccc cgagcgccac cggcaggcc gaggcatcc ggcgtctgt cgaccggacc 3960
ggcattcgacc ctgcattcgat tggctacgtt gaggccatg gcaccggaa ctgtgggt 4020
gaccctgtcg aggctccgc gctgaggcga gcttcggc ctttcacccga cggcggcggg 4080
tactgcccgc tgggtcggt gaagtcaacg ctggccatc tggacacagt ggctggactg 4140
gctgggtc tcaagacggc gctgaggcc cggcaggcc aagtccctc gacgcctcat 4200
gtgaccagg tgaatccaa gctcgaggc acggattcgc ctttcgtcat cggcggaccgt 4260
ttggcgcgtt ggccgtccct gccgggaccg aggccggcgg cctgtggcgt gtcggccctt 4320
ggcgggacga atacccacgc catttcgaa cactacccgc ggcactcccg cccacgggg 4380
aggagccacg ggtcaacgc agtccgtcg tgggtccat tctccgcgc caccctggag 4440
gcgttgaagg acaaccccg cgcgtgtc gacttcgttgg aggaccggc gtccgggg 4500
gtggcgctcg cggacatcac ctacacgtt caggccggc ggttgcgtat gcttgcggg 4560
atgggtgttgc tgcgtcgac ggcgcacgaa ttgggtggagg gacttgcggc aggcatcgcc 4620
acggtggcgc gtggccacgtt gggaaacgggtt gtcgatcgatc caccacgtt ggttgcggat 4680
gctcggcag ttggggaggc gtggggcgcg ggcaacttgcg ttgactgggat ttcgtgcac 4740
ggtgacgttgc agccggcccg tgcacgttgc cccacgttgc agtttgcgaa ggagcgctac 4800
gggttgcgc cccgcacac tgcgttgcgaa ttcccaaga gtcacatccatc cgcgggtgtc 4860
ccgcgttgc ttcggaccgtt gcaacccgtt tctgttgcgc ttggggcttgc aggttgcctt 4920
ctccggcacc tgggtgttgc tgcgttgcgc ttgtatgcgc ttggggcttgc aggttgcctt 4980
gcgcgttgcgca gacgttgcgc ggcacaggccg atcgaatgtt ctttcgttgc cagccaaatgt 5040
gcgcggctgg acggccgggtt catggccatc gcttcggcgg ttttcgttgc acgttgcgg 5100
ctgtgttgcg acgggttgc ctttcgttgc acatttcgttgc ttgttgcgttgc agggaggccg 5160
gtatgttgcg cacttgcgttgc ctttcgttgc acgttgcgttgc ttgttgcgttgc ggagaatccg 5220
ttggccggg ggcacgttgc cccgttgcgttgc ggaacccgttgc cccgttgcgttgc ttgttgcgttgc 5280
tttctgggttgc agtccgcgcg cggccgttgc acgttgcgttgc cccgttgcgttgc ttgttgcgttgc 5340
cttttcgttgc ttgttgcgttgc acgttgcgttgc cccgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5400
cgcaagacg gcttgcgttgc gatgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5460
ggcgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5520
ggcgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5580
caacccgttgc acgttgcgttgc ctttcgttgc acgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5640
gggttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5700
gaaatgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5760
cgcaacccgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5820
gggttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5880
ggcgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 5940
gagaacccgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6000
atggccggccg tggggggggcc acggggacttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6060
tccctgggttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6120
ttttccgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6180
aagggttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6240
atcgatccgc gcaacccgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6300
aaccacccgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6360
acgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6420
gtggccggccac ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6480
gatgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6540
cggttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6600
tgcgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6660
aaggacccgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6720
gctgttgcgttgc acggccgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc ttgttgcgttgc 6780

caggagcgct tggccctgca gagctgcctgg gaggtccctgg aggacgcggg gtacacccgg 6840
gacagcctgg cccagcgctt tggccagcgcg gtggcggtt tcgcggaaat cacgaagacg 6900
ggctacgaac tctacggcgc ggagctggaa ggacgagatg cctcggtccg gcctatacg 6960
tcgttgcgt ctgtgcctaa cccgtcgatc tatctgcctcg acctgaaggg gccgagcatg 7020
cccggtggaca ccatgtgcgc ggcctcgatc acagccgtcc acatggctcg cgaggcgctg 7080
caacgaggcg cctgcgtcat ggcctcgatc ggtggagtga atctctacgt ccacccgtcg 7140
actacgtca gcctgtcccg gcagcagatg ctgtcgac 7178

<210> 3
<211> 785
<212> DNA
<213> Myxococcus xanthus

<400> 3
Met Lys Val Val Asn Lys Leu Leu Glu Lys Leu Pro Asp Val Val Ala
1 5 10 15

Gly Lys Val Pro Asp Val Lys Leu Gln Asp Gln Asp Ile Lys Val Pro
20 25 30

Leu Ala Gln Gly Thr Phe Thr Glu Glu Lys Ile Leu Pro Pro Lys Leu
35 40 45

Ala Met His Gly Phe Thr Leu Ser Phe Glu Ala Thr Gly Glu Ala Ser
50 55 60

Ile Arg Asn Phe Asn Ser Leu Gly Asp Val Asp Glu Asn Gly Ile Ile
65 70 75 80

Gly Glu Pro Ser Pro Glu Ser Ala Glu Pro Gly Pro Arg Pro Gln Leu
85 90 95

Leu Leu Gly Ser Asp Ile Gly Trp Met Arg Tyr Gln Val Ser Ala Arg
100 105 110

Val Lys Ala Ala Val Ser Ala Ser Leu Ser Phe Leu Ala Ser Glu Asn
115 120 125

Gln Thr Glu Leu Ser Val Thr Leu Ser Asp Tyr Arg Ala His Pro Leu
130 135 140

Gly Gln Asn Met Arg Glu Ala Val Arg Ser Asp Leu Ser Glu Leu Arg
145 150 155 160

Leu Met Gln Ala Thr Asp Leu Ala Lys Leu Thr Thr Gly Asp Ala Val
165 170 175

Ala Trp His Val Arg Gly Ala Leu His Thr Arg Leu Glu Leu Asn Trp
180 185 190

Ala Asp Ile Phe Pro Thr Asn Leu Asn Arg Leu Gly Phe Leu Arg Gly
195 200 205

Asn Glu Leu Leu Ala Leu Lys Thr Ser Ala Lys Ala Gly Leu Ser Ala
210 215 220

Arg Val Ser Leu Thr Asp Asp Tyr Gln Leu Ser Phe Ser Arg Pro Arg
225 230 235 240

Ala Gly Arg Ile Gln Val Ala Val Arg Lys Val Lys Ser His Glu Gln
245 250 255

Ala Leu Ser Ala Gly Leu Gly Ile Thr Val Glu Leu Leu Asp Pro Ala
260 265 270

Thr Val Lys Ala Gln Leu Gly Gln Leu Leu Glu Ala Leu Leu Gly Pro
275 280 285

Val Leu Arg Asp Leu Val Lys Lys Gly Thr Thr Ala Val Glu Ile Met
290 295 300

Asp Gly Leu Val Asp Lys Ala Ser Lys Ala Lys Leu Asp Asp Asn Gln
305 310 315 320

Lys Lys Val Leu Gly Leu Val Leu Glu Arg Leu Gly Ile Asp Pro Gln
325 330 335

Leu Ala Asp Pro Ala Asn Leu Pro Gln Ala Trp Ala Asp Phe Lys Ala
340 345 350

Arg Val Ala Glu Ser Leu Glu Asn Ala Val Arg Thr Gln Val Ala Glu
355 360 365

Gly Phe Glu Tyr Glu Tyr Leu Arg Leu Ser Glu Thr Ser Thr Leu Leu
370 375 380

Glu Val Val Val Glu Asp Val Thr Ala Met Arg Phe His Glu Ser Leu
385 390 395 400

Leu Lys Gly Asn Leu Val Glu Leu Leu Lys Trp Met Lys Ser Leu Pro
405 410 415

Ala Gln Gln Ser Glu Phe Glu Leu Arg Asn Tyr Leu His Ala Thr Thr
420 425 430

Leu Thr Arg Gln Gln Ala Ile Gly Phe Ser Leu Gly Leu Gly Ser Phe
435 440 445

Glu Leu Leu Lys Ala Lys Asn Val Ser Lys Gln Ser Trp Val Thr Gln
450 455 460

Glu Asn Phe Gln Gly Ala Arg Arg Met Ala Phe Leu Gly Arg Arg Gly
465 470 475 480

Tyr Glu Asp Lys Leu Leu Gly Thr Arg Gly Gln Trp Val Val Asp Leu
485 490 495

Lys Ala Asp Met Thr Arg Phe Ser Pro Thr Pro Val Ala Ser Asp Phe
500 505 510

Gly Tyr Gly Leu His Leu Met Leu Trp Gly Arg Gln Lys Lys Leu Ser
515 520 525

Arg Lys Asp Leu Gln Gln Ala Val Asp Asp Ala Val Val Trp Gly Val
530 535 540

Leu Asp Ala Lys Asp Ala Ala Thr Val Ile Ser Thr Met Gln Glu Asp
545 550 555 560

Met Gly Lys His Pro Ile Glu Thr Arg Leu Glu Leu Lys Met Ala Asp
565 570 575

Asp Ser Phe Arg Ala Leu Val Pro Arg Ile Glu Thr Leu Glu Leu Ser
580 585 590

Arg Phe Ser Arg Ala Leu Ala Arg Ala Leu Pro Trp Ser Glu Gln Leu
595 600 605

Pro Arg Ala Ser Ala Glu Phe Arg Arg Ala Val Tyr Ala Pro Ile Trp
610 615 620

Glu Ala Tyr Leu Arg Glu Val Gln Glu Gln Gly Ser Leu Met Leu Asn
625 630 635 640

Asp Leu Ser Pro Ser Arg Ala Ala Gln Ile Ala Lys Trp Tyr Phe Gln
645 650 655

Lys Asp Pro Thr Val Arg Asp Leu Gly Lys Asp Leu Gln Leu Ne Glu
660 665 670

Ser Glu Trp Arg Pro Gly Gly Asn Phe Ser Phe Ala Glu Val Ile
675 680 685

Ser Lys Asn Pro Asn Thr Leu Met Arg Cys Arg Asn Phe Val Ser Gly
690 695 700

Met Val Arg Leu Arg Arg Ala Ile Asp Glu Arg Lys Ala Pro Asp Glu
705 710 715 720

Leu Arg Thr Val Phe Gly Glu Leu Glu Gly Met Trp Thr Thr Gly Phe
725 730 735

His Leu Arg Ala Ala Gly Ser Leu Leu Ser Asp Leu Ala Gln Ser Thr
740 745 750

Pro Leu Gly Leu Ala Gly Val Glu Arg Thr Leu Thr Val Arg Val Ala
755 760 765

Asp Ser Glu Glu Gln Leu Val Phe Ser Thr Ala Arg Ser Thr Gly Ala
770 775 780

Ala
785

<210> 4
<211> 529
<212> DNA
<213> Myxococcus xanthus

<400> 4
Met Pro Ser Gly Cys Tyr Gly Ala Ala Ser Ala Phe Val Leu Pro Pro
1 5 10 15

Leu Pro Ala Met Pro Gln Ala Pro Ser Asp Val Ser Gln Val Leu Leu
20 25 30

Pro Phe Gly Gly Leu Val Gly Arg Glu Val Asp Leu Asp Ala Phe Leu
35 40 45

Gln Thr Leu Met Asp Arg Ile Ala Ile Thr Leu Gln Ala Asp Arg Gly
50 55 60

Thr Leu Trp Leu Leu Asp Pro Ala Arg Arg Glu Leu Phe Ser Arg Ala
65 70 75 80

Ala His Leu Pro Glu Val Ser Gln Ile Arg Val Lys Leu Gly Gln Gly
85 90 95

Val Ala Gly Thr Val Ala Lys Ala Gly His Ala Ile Asn Val Pro Asp
100 105 110

Pro Arg Gly Glu Gln Arg Phe Phe Ala Asp Ile Asp Arg Met Thr Gly
115 120 125

Tyr Arg Thr Thr Ser Leu Leu Ala Val Pro Leu Arg Asp Gly Asp Gly
130 135 140

Ala Leu Tyr Gly Val Leu Gln Val Leu Asn Arg Arg Gly Glu Asp Arg
145 150 155 160

Phe Thr Asp Glu Asp Thr Gln Arg Leu Thr Ala Ile Ala Ser Gln Val
165 170 175

Ser Thr Ala Leu Gln Ser Thr Ser Leu Tyr Gln Glu Leu Gln Arg Ala
180 185 190

Lys Glu Gln Pro Gln Val Pro Val Gly Tyr Phe Asn Arg Ile Ile
195 200 205

Gly Glu Ser Pro Gln Leu Gln Ala Ile Tyr Arg Leu Val Arg Lys Ala
210 215 220

Ala Pro Thr Asp Ala Thr Val Leu Leu Arg Gly Glu Ser Gly Ser Gly
225 230 235 240

Lys Glu Leu Phe Ala Arg Ala Val His Val Asn Gly Pro Arg Arg Asp
245 250 255

Gln Pro Phe Ile Lys Val Asp Cys Ala Ala Leu Pro Ala Thr Leu Ile
260 265 270

Glu Asn Glu Leu Phe Gly His Glu Arg Gly Ala Phe Thr Gly Ala Asp
275 280 285

His Arg Val Pro Gly Lys Phe Glu Ala Ala Ser Gly Gly Thr Val Phe
290 295 300

Ile Asp Glu Ile Gly Glu Leu Pro Leu Pro Val Gln Gly Lys Leu Leu
305 310 315 320

Arg Val Ile Gln Asp Arg Glu Phe Glu Arg Val Gly Gly Thr Gln Ala
325 330 335

Val Lys Val Asp Val Arg Ile Val Ala Ala Thr His Arg Asp Leu Ala
340 345 350

Arg Met Val Ala Glu Gly Arg Phe Arg Glu Asp Leu Tyr Tyr Arg Ile
355 360 365

Lys Val Val Glu Val Val Leu Pro Pro Leu Arg Glu Arg Gly Ala Glu
370 375 380

Asp Ile Glu Arg Leu Ala Arg His Phe Val Ala Ala Val Ala Arg Arg
385 390 395 400

His Arg Leu Thr Pro Pro Arg Leu Ser Ala Ala Ala Val Glu Arg Leu
405 410 415

Lys Arg Tyr Arg Trp Pro Gly Asn Val Arg Glu Leu Glu Asn Cys Ile
420 425 430

Glu Ser Ala Val Val Leu Cys Glu Gly Glu Ile Leu Glu Glu His Leu
435 440 445

Pro Leu Pro Asp Val Asp Arg Ala Ala Leu Pro Pro Pro Ala Ala Ala
450 455 460

Gln Gly Val Asn Ala Pro Thr Ala Pro Ala Pro Leu Asp Ala Gly Leu
465 470 475 480

Leu Pro Leu Ala Glu Val Glu Arg Arg His Ile Leu Arg Val Leu Asp
485 490 495

Ala Val Lys Gly Asn Arg Thr Ala Ala Arg Val Leu Ala Ile Gly
500 505 510

Arg Asn Thr Leu Ala Arg Lys Leu Lys Glu Tyr Gly Leu Gly Asp Glu
515 520 525

Pro

<210> 5
<211> 292
<212> DNA
<213> *Myxococcus xanthus*

<400> 5

Met Arg Ala Ser Gln Ala Glu Ala Pro His Ser Arg Arg Leu Thr Met
1 5 10 15

Glu Val Arg Phe His Gly Val Arg Gly Ser Ile Ala Val Ser Gly Ser
20 25 30

Arg Ile Gly Gly Asn Thr Ala Cys Val Glu Val Thr Ser Gln Gly His
35 40 45

Arg Leu Ile Leu Asp Ala Gly Thr Gly Ile Arg Ala Leu Gly Glu Ile
50 55 60

Met Met Arg Glu Gly Ala Pro Gln Glu Ala Thr Leu Phe Phe Ser His
65 70 75 80

Leu His Trp Asp His Val Gln Gly Phe Pro Phe Phe Thr Pro Ala Trp
85 90 95

Leu Pro Thr Ser Glu Leu Thr Leu Tyr Gly Pro Gly Ala Asn Gly Ala
100 105 110

Gln Ala Leu Gln Ser Glu Leu Ala Ala Gln Met Gln Pro Leu His Phe
115 120 125

Pro Val Pro Leu Ser Thr Met Arg Ser Arg Met Asp Phe Arg Ser Ala
130 135 140

Leu His Ala Arg Pro Val Glu Val Gly Pro Phe Arg Val Thr Pro Ile
145 150 155 160

Asp Val Pro His Pro Gln Gly Cys Leu Ala Tyr Arg Leu Glu Ala Asp
165 170 175

Gly His Ser Phe Val Tyr Ala Thr Asp Val Glu Val Arg Val Gln Glu
180 185 190

Leu Ala Pro Glu Val Gly Arg Leu Phe Glu Gly Ala Asp Val Leu Cys
195 200 205

Leu Asp Ala Gln Tyr Thr Pro Asp Glu Tyr Glu Gly Arg Lys Gly Val
210 215 220

Ala Lys Lys Gly Trp Gly His Ser Thr Met Met Asp Ala Ala Gly Val

225 230 235 240
Ala Gly Leu Val Gly Ala Arg Arg Leu Cys Leu Phe His His Asp Pro
245 250 255
Ala His Gly Asp Asp Met Leu Glu Asp Met Ala Glu Gln Ala Arg Ala
260 265 270
Leu Phe Pro Val Cys Glu Pro Ala Arg Glu Gly Gln Arg Leu Val Leu
275 280 285
Gly Arg Ala Ala
290

<210> 6
<211> 168
<212> DNA
<213> *Myxococcus xanthus*

<400> 6
Met Pro Gly Pro Arg Cys Ala Glu Asn Asp Trp Val Ala Leu Leu Val
1 5 10 15
Arg Val Asn His Glu Lys Val Ala Ala Gln Leu Gly Lys His Gly
20 25 30
Tyr Glu Phe Phe Leu Pro Thr Tyr Thr Pro Pro Lys Ser Ser Gly Val
35 40 45
Lys Ala Lys Leu Pro Leu Phe Pro Gly Tyr Leu Phe Cys Arg Tyr Gln
50 55 60
Pro Leu Asn Pro Tyr Arg Ile Val Arg Ala Pro Gly Val Ile Arg Leu
65 70 75 80
Leu Gly Gly Asp Ala Gly Pro Glu Ala Val Pro Ala Gln Glu Leu Glu
85 90 95
Ala Ile Arg Arg Val Ala Asp Ser Gly Val Ser Ser Asn Pro Cys Asp
100 105 110
Tyr Leu Arg Val Gly Gln Arg Val Arg Ile Ile Glu Gly Pro Leu Thr
115 120 125
Gly Leu Glu Gly Ser Leu Val Thr Ser Lys Ser Gln Leu Arg Phe Ile
130 135 140
Val Ser Val Gly Leu Leu Gln Arg Ser Val Ser Val Glu Val Ser Ala
145 150 155 160
Glu Gln Leu Glu Pro Ile Thr Asp
165

<210> 7
<211> 79
<212> DNA
<213> *Myxococcus xanthus*

<400> 7
Met Asp Lys Arg Ile Ile Phe Asp Ile Val Thr Ser Ser Val Arg Glu
1 5 10 15

Val Val Pro Glu Leu Glu Ser His Pro Phe Glu Pro Glu Asp Asp Leu
20 25 30

Val Gly Leu Gly Ala Asn Ser Leu Asp Arg Ala Glu Ile Val Asn Leu
35 40 45

Thr Leu Glu Lys Leu Ala Leu Asn Ile Pro Arg Val Glu Leu Ile Asp
50 55 60

Ala Lys Thr Ile Gly Gly Leu Val Asp Val Leu His Ala Arg Leu
65 70 75

<210> 8
<211> 420
<212> DNA
<213> *Myxococcus xanthus*

<400> 8
Met Gly Pro Val Gly Ile Glu Ala Met Asn Ala Tyr Cys Gly Ile Ala
1 5 10 15

Arg Leu Asp Val Leu Gln Leu Ala Thr His Arg Gly Leu Asp Thr Ser
20 25 30

Arg Phe Ala Asn Leu Leu Met Glu Glu Lys Thr Val Pro Leu Pro Tyr
35 40 45

Glu Asp Pro Val Thr Tyr Gly Val Asn Ala Ala Arg Pro Ile Leu Asp
50 55 60

Gln Leu Thr Ala Ala Glu Arg Asp Ser Ile Glu Leu Leu Val Ala Cys
65 70 75 80

Thr Glu Ser Ser Phe Asp Phe Gly Lys Ala Met Ser Thr Tyr Leu His
85 90 95

Gln His Leu Gly Leu Ser Arg Asn Cys Arg Leu Ile Glu Leu Lys Ser
100 105 110

Ala Cys Tyr Ser Gly Val Ala Gly Leu Gln Met Ala Val Asn Phe Ile
115 120 125

Leu Ser Gly Val Ser Pro Gly Ala Lys Ala Leu Val Val Ala Ser Asp

130	135	140
Leu Ser Arg Phe Ser Ile Ala Glu Gly Gly Asp Ala Ser Thr Glu Asp		
145	150	155
Trp Ser Phe Ala Glu Pro Ser Ser Gly Ala Gly Ala Val Ala Met Leu		
165	170	175
Val Ser Asp Thr Pro Arg Val Phe Arg Val Asp Val Gly Ala Asn Gly		
180	185	190
Tyr Tyr Gly Tyr Glu Val Met Asp Thr Cys Arg Pro Val Ala Asp Ser		
195	200	205
Glu Ala Gly Asp Ala Asp Leu Ser Leu Leu Ser Tyr Leu Asp Cys Cys		
210	215	220
Glu Asn Ala Phe Arg Glu Tyr Thr Arg Arg Val Pro Ala Ala Asn Tyr		
225	230	235
Ala Glu Ser Phe Gly Tyr Leu Ala Phe His Thr Pro Phe Gly Gly Met		
245	250	255
Val Lys Gly Ala His Arg Thr Met Met Arg Lys Phe Ser Gly Lys Asn		
260	265	270
Arg Gly Asp Ile Glu Ala Asp Phe Gln Arg Arg Val Ala Pro Gly Leu		
275	280	285
Thr Tyr Cys Gln Arg Val Gly Asn Ile Met Gly Ala Thr Met Ala Leu		
290	295	300
Ser Leu Leu Gly Thr Ile Asp His Gly Asp Phe Ala Thr Ala Lys Arg		
305	310	315
Ile Gly Cys Phe Ser Tyr Gly Ser Gly Cys Ser Ser Glu Phe Phe Ser		
325	330	335
Gly Val Val Thr Glu Glu Gly Gln Gln Arg Gln Arg Ala Leu Gly Leu		
340	345	350
Gly Glu Ala Leu Gly Arg Arg Gln Gln Leu Ser Met Pro Asp Tyr Asp		
355	360	365
Ala Leu Leu Lys Gly Asn Gly Leu Val Arg Phe Gly Thr Arg Asn Ala		
370	375	380
Glu Leu Asp Phe Gly Val Val Gly Ser Ile Arg Pro Gly Gly Trp Gly		
385	390	395
Arg Pro Leu Leu Phe Leu Ser Ala Ile Arg Asp Phe His Arg Asp Tyr		
405	410	415
Gln Trp Ile Ser		

420
<210> 9
<211> 325
<212> DNA
<213> *Myxococcus xanthus*

<400> 9

Met Ser Ser Val Ala Thr Ala Val Pro Leu Thr Ala Arg Asp Ser Ala
1 5 10 15

Val Ser Arg Arg Leu Arg Ile Thr Pro Ser Met Cys Gly Gln Thr Ser
20 25 30

Leu Phe Ala Gly Gln Ile Gly Asp Trp Ala Trp Asp Thr Val Ser Arg
35 40 45

Leu Cys Gly Thr Asp Val Leu Thr Ala Thr Asn Ala Ser Gly Ala Pro
50 55 60

Thr Tyr Leu Ala Phe Tyr Tyr Phe Arg Ile Arg Gly Thr Pro Ala Leu
65 70 75 80

His Pro Gly Ala Leu Arg Phe Gly Asp Thr Leu Asp Val Thr Ser Lys
85 90 95

Ala Tyr Asn Phe Gly Ser Glu Ser Val Leu Thr Val His Arg Ile Cys
100 105 110

Lys Thr Ala Glu Gly Ala Pro Glu Ala Asp Ala Phe Gly His Glu
115 120 125

Glu Leu Tyr Glu Gln Pro Gln Pro Gly Arg Ile Tyr Ala Glu Thr Phe
130 135 140

Asn Arg Trp Ile Thr Arg Ser Asp Gly Lys Ser Asn Glu Ser Leu Ile
145 150 155 160

Lys Ser Ser Pro Val Gly Phe Gln Tyr Ala His Leu Pro Leu Leu Pro
165 170 175

Asp Glu Tyr Ser Pro Arg Arg Ala Tyr Gly Asp Ala Arg Ala Arg Gly
180 185 190

Thr Phe His Asp Val Asp Ser Ala Glu Tyr Arg Leu Thr Val Asp Arg
195 200 205

Phe Pro Leu Arg Tyr Ala Val Asp Val Ile Arg Asp Val Asn Gly Val
210 215 220

Gly Leu Ile Tyr Phe Ala Ser Tyr Phe Ser Met Val Asp Trp Ala Ile
225 230 235 240

Trp Gln Leu Ala Arg His Gln Gly Arg Ser Glu Gln Ala Phe Leu Ser
245 250 255

Arg Val Val Leu Asp Gln Gln Leu Cys Phe Leu Gly Asn Ala Ala Leu
260 265 270

Asp Thr Thr Phe Asp Ile Asp Val Gln His Trp Glu Arg Val Gly Gly
275 280 285

Gly Glu Glu Leu Phe Asn Val Lys Met Arg Glu Gly Ala Gln Gly Arg
290 295 300

Asp Ile Ala Val Ala Thr Val Lys Val Arg Phe Asp Ala Ala Ser Glu
305 310 315 320

Gly Gly Arg Arg Gly
325

<210> 10

<211> 83

<212> DNA

<213> Myxococcus xanthus

<400> 10

Met Thr Asp Glu Gln Ile Arg Gly Val Val His Gln Ser Ile Val Arg
1 5 10 15

Val Leu Pro Arg Val Arg Ser Asn Glu Ile Ala Gly His Leu Asn Leu
20 25 30

Arg Glu Leu Gly Ala Asp Ser Val Asp Arg Val Glu Ile Leu Thr Ser
35 40 45

Ile Leu Asp Ser Leu Arg Leu Gln Lys Thr Pro Leu Ala Lys Phe Ala
50 55 60

Asp Ile Arg Asn Ile Asp Ala Leu Val Ala Phe Leu Ala Gly Glu Val
65 70 75 80

Ala Gly Gly

<210> 11

<211> 374

<212> DNA

<213> Myxococcus xanthus

<400> 11

Met Met Gln Glu Arg Gly Val Ala Leu Pro Phe Glu Asp Pro Val Thr
1 5 10 15

Asn Ala Val Asn Ala Ala Arg Pro Ile Leu Asp Ala Met Ser Pro Glu

20 25 30
Ala Arg Glu Arg Ile Glu Leu Leu Val Thr Ser Ser Glu Ser Gly Val
35 40 45
Asp Phe Ser Lys Ser Ile Ser Ser Tyr Ala His Glu His Leu Gly Leu
50 55 60
Ser Arg His Cys Arg Phe Leu Glu Val Lys Gln Ala Cys Tyr Ala Ala
65 70 75 80
Thr Gly Ala Leu Gln Leu Ala Leu Gly Tyr Ile Ala Ser Gly Val Ser
85 90 95
Pro Gly Ala Lys Ala Leu Val Ile Ala Thr Asp Val Thr Leu Val Asp
100 105 110
Glu Ser Gly Leu Tyr Ser Glu Pro Ala Met Gly Thr Gly Gly Val Ala
115 120 125
Val Leu Leu Gly Asp Glu Pro Arg Val Met Lys Met Asp Leu Gly Ala
130 135 140
Phe Gly Asn Tyr Ser Tyr Asp Val Phe Asp Thr Ala Arg Pro Ser Pro
145 150 155 160
Glu Ile Asp Ile Gly Asp Val Asp Arg Ser Leu Phe Thr Tyr Leu Asp
165 170 175
Cys Leu Lys His Ser Phe Ala Ala Tyr Gly Arg Arg Val Asp Gly Val
180 185 190
Asp Phe Val Ser Thr Phe Asp Tyr Leu Ala Met His Thr Pro Phe Ala
195 200 205
Gly Leu Val Lys Ala Gly His Arg Lys Met Met Arg Glu Leu Thr Pro
210 215 220
Cys Asp Val Asp Glu Ile Glu Ala Asp Phe Gly Arg Arg Val Lys Pro
225 230 235 240
Ser Leu Gln Tyr Pro Ser Leu Val Gly Asn Leu Cys Ser Gly Ser Val
245 250 255
Tyr Leu Ser Leu Cys Ser Ile Ile Asp Thr Ile Lys Pro Glu Arg Ser
260 265 270
Ala Arg Val Gly Met Phe Ser Tyr Gly Ser Gly Cys Ser Ser Glu Phe
275 280 285
Phe Ser Gly Val Ile Gly Pro Glu Ser Val Ser Ala Leu Ala Gly Leu
290 295 300
Asp Ile Gly Gly His Leu Arg Gly Arg Arg Gln Leu Thr Phe Asp Gln

305 310 315 320
Tyr Val Glu Leu Leu Lys Glu Asn Leu Arg Cys Leu Val Pro Thr Lys
325 330 335
Asn Arg Asp Val Asp Val Glu Arg Tyr Leu Pro Leu Val Thr Arg Thr
340 345 350
Ala Ser Arg Pro Arg Met Leu Ala Leu Arg Arg Val Val Asp Tyr His
355 360 365
Arg Gln Tyr Glu Trp Val
370

<210> 12
<211> 171
<212> DNA
<213> *Myxococcus xanthus*

<400> 12
Met Asn Thr Pro Ser Leu Thr Asn Trp Pro Ala Arg Leu Gly Tyr Leu
1 5 10 15
Leu Ala Val Gly Gly Ala Trp Phe Ala Ala Asp Gln Val Thr Lys Gln
20 25 30
Met Ala Arg Asp Gly Ala Lys Arg Pro Val Ala Val Phe Asp Ser Trp
35 40 45
Trp His Phe His Tyr Val Glu Asn Arg Ala Gly Ala Phe Gly Leu Phe
50 55 60
Ser Ser Phe Gly Glu Glu Trp Arg Met Pro Phe Phe Tyr Val Val Gly
65 70 75 80
Ala Ile Cys Ile Val Leu Leu Ile Gly Tyr Tyr Phe Tyr Thr Pro Pro
85 90 95
Thr Met Lys Leu Gln Arg Trp Ser Leu Ala Thr Met Ile Gly Gly Ala
100 105 110
Leu Gly Asn Tyr Val Asp Arg Val Arg Leu Arg Tyr Val Val Asp Phe
115 120 125
Val Ser Trp His Val Gly Asp Arg Phe Tyr Trp Pro Ser Phe Asn Ile
130 135 140
Ala Asp Thr Ala Val Val Val Gly Ala Ala Leu Met Ile Leu Glu Ser
145 150 155 160
Phe Arg Glu Pro Arg Gln Gln Leu Ser Pro Gly
165 170

<210> 13
<211> 475
<212> DNA
<213> *Myxococcus xanthus*

<400> 13

Met Gly Thr Ser Glu Pro Val Glu Pro Asp His Ala Leu Ser Lys Pro
1 5 10 15

Pro Pro Val Ala Pro Val Gly Ala Gln Ala Leu Pro Arg Gly Pro Ala
20 25 30

Met Pro Gly Ile Ala Gln Leu Met Met Leu Phe Leu Arg Pro Thr Glu
35 40 45

Phe Leu Asp Arg Cys Ala Ala Arg Tyr Gly Asp Thr Phe Thr Leu Lys
50 55 60

Ile Pro Gly Thr Pro Pro Phe Ile Gln Thr Ser Asp Pro Ala Leu Ile
65 70 75 80

Glu Val Ile Phe Lys Gly Asp Pro Asp Leu Phe Leu Gly Gly Lys Ala
85 90 95

Asn Asn Gly Leu Lys Pro Val Val Gly Glu Asn Ser Leu Leu Val Leu
100 105 110

Asp Gly Lys Arg His Arg Arg Asp Arg Lys Leu Ile Met Pro Thr Phe
115 120 125

Leu Gly Glu Arg Met His Ala Tyr Gly Ser Val Ile Arg Asp Ile Val
130 135 140

Asn Ala Ala Leu Asp Arg Trp Pro Val Gly Lys Pro Phe Ala Val His
145 150 155 160

Glu Glu Thr Gln Gln Ile Met Leu Glu Val Ile Leu Arg Val Ile Phe
165 170 175

Gly Leu Glu Asp Ala Arg Thr Ile Ala Gln Phe Arg His His Val His
180 185 190

Gln Val Leu Lys Leu Ala Leu Phe Leu Phe Pro Asn Gly Glu Gly Lys
195 200 205

Pro Ala Ala Glu Gly Phe Ala Arg Ala Val Gly Lys Ala Phe Pro Ser
210 215 220

Leu Asp Val Phe Ala Ser Leu Lys Ala Ile Asp Asp Ile Ile Tyr Gln
225 230 235 240

Glu Ile Gln Asp Arg Arg Ser Gln Asp Ile Ser Gly Arg Gln Asp Val
245 250 255

Leu Ser Leu Met Met Gln Ser His Tyr Asp Asp Gly Ser Val Met Thr
260 265 270

Pro Gln Glu Leu Arg Asp Glu Leu Met Thr Leu Leu Met Ala Gly His
275 280 285

Glu Thr Ser Ala Thr Ile Ala Ala Trp Cys Val Tyr His Leu Cys Arg
290 295 300

His Pro Asp Ala Met Gly Lys Leu Arg Glu Glu Ile Ala Ala His Thr
305 310 315 320

Val Asp Gly Val Leu Pro Leu Ala Lys Ile Asn Glu Leu Lys Phe Leu
325 330 335

Asp Ala Val Val Lys Glu Thr Met Arg Ile Thr Pro Val Phe Ser Leu
340 345 350

Val Ala Arg Val Leu Lys Glu Pro Gln Thr Ile Gly Gly Thr Thr Tyr
355 360 365

Pro Ala Asn Val Val Leu Ser Pro Asn Ile Tyr Gly Thr His His Arg
370 375 380

Ala Asp Leu Trp Gly Asp Pro Lys Val Phe Arg Pro Glu Arg Phe Leu
385 390 395 400

Glu Glu Arg Val Asn Pro Phe His Tyr Phe Pro Phe Gly Gly Ile
405 410 415

Arg Lys Cys Ile Gly Thr Ser Phe Ala Tyr Tyr Glu Met Lys Ile Phe
420 425 430

Val Ser Glu Thr Val Arg Arg Met Arg Phe Asp Thr Arg Pro Gly Tyr
435 440 445

His Ala Lys Val Val Arg Arg Ser Asn Thr Leu Ala Pro Ser Gln Gly
450 455 460

Val Pro Ile Ile Val Glu Ser Arg Leu Pro Ser
465 470 475

<210> 14

<211> 318

<212> DNA

<213> *Myxococcus xanthus*

<400> 14

Met Val Asp Ser Val Ser Lys Gln Ala Arg Arg Lys Val Phe Leu Phe
1 5 10 15

Ser Gly Gln Gly Thr Gln Ser Tyr Phe Met Ala Lys Glu Leu Phe Asp

20 25 30
Thr Gln Thr Gly Rhe Lys Arg Gln Leu Leu Glu Leu Asp Glu Gln Phe
35 40 45
Lys Gln Arg Leu Gly His Ser Ile Leu Glu Arg Ile Tyr Asp Ala Arg
50 55 60
Ala Ala Arg Leu Asp Pro Leu Asp Asp Val Leu Val Ser Phe Pro Ala
65 70 75 80
Ile Phe Met Ile Glu His Ala Leu Ala Arg Leu Leu Ile Asp Arg Gly
85 90 95
Ile Gln Pro Asp Ala Val Val Gly Ala Ser Met Gly Glu Val Ala Ala
100 105 110
Ala Ala Ile Ala Gly Ala Ile Ser Val Asp Ala Ala Val Ala Leu Val
115 120 125
Ala Ala Gln Ala Gln Leu Phe Ala Arg Thr Ala Pro Arg Gly Gly Met
130 135 140
Leu Ala Val Leu His Glu Leu Glu Ala Cys Arg Gly Phe Thr Ser Val
145 150 155 160
Ala Arg Asp Gly Glu Val Ala Ala Ile Asn Tyr Pro Ser Asn Phe Val
165 170 175
Leu Ala Ala Asp Glu Ala Gly Leu Gly Arg Ile Gln Gln Glu Leu Ser
180 185 190
Gln Arg Ser Val Ala Phe His Arg Leu Pro Val Arg Tyr Pro Phe His
195 200 205
Ser Ser His Leu Asp Pro Leu Arg Glu Glu Tyr Arg Ser Arg Val Arg
210 215 220
Ala Asp Ser Leu Thr Trp Pro Arg Ile Pro Met Tyr Ser Cys Thr Thr
225 230 235 240
Ala Asn Arg Val His Asp Leu Arg Ser Asp His Phe Trp Asn Val Val
245 250 255
Arg Ala Pro Ile Gln Leu Tyr Asp Thr Val Leu Gln Leu Glu Gly Gln
260 265 270
Gly Gly Cys Asp Phe Ile Asp Val Gly Pro Ala Ala Ser Phe Ala Thr
275 280 285
Ile Ile Lys Arg Ile Leu Ala Arg Asp Ser Thr Ser Arg Leu Phe Pro
290 295 300
Leu Leu Ser Pro Ser Pro Ala Ser Thr Gly Ser Ser Met Gly

B
Cmt

305

310

315

<210> 15

<211> 330

<212> DNA

<213> *Myxococcus xanthus*

<400> 15

Met Thr Glu Ala Pro Ala Pro Arg Ala Pro Ala Gln Val Pro Pro Pro
 1 5 10 15

Pro Ser Ser Pro Trp Ala Leu His Thr Arg Gly Ala Ala Ser Ala Pro
 20 25 30

Val Asn Ala Arg Lys Ala Ala Leu Phe Pro Gly Gln Gly Ser Gln Glu
 35 40 45

Arg Gly Met Gly Ala Ala Leu Phe Asp Glu Phe Pro Asp Leu Thr Asp
 50 55 60

Ile Ala Asp Ala Ile Leu Gly Tyr Ser Ile Lys Arg Leu Cys Leu Glu
 65 70 75 80

Asp Pro Gly Lys Glu Leu Ala Gln Thr Gln Phe Thr Gln Pro Ala Leu
 85 90 95

Tyr Val Val Asn Ala Leu Ser Tyr Leu Lys Arg Leu Arg Glu Gly Ala
 100 105 110

Glu Gln Pro Ala Phe Val Ala Gly His Ser Leu Gly Glu Tyr Asn Ala
 115 120 125

Leu Leu Val Ala Gly Ala Phe Asp Phe Glu Thr Gly Leu Arg Leu Val
 130 135 140

Lys Arg Arg Gly Glu Leu Met Ser Gly Ala Ser Gly Gly Thr Met Ala
 145 150 155 160

Ala Val Val Gly Cys Asp Ala Val Ala Val Glu Gln Val Leu Arg Asp
 165 170 175

Arg Gln Leu Thr Ser Leu Asp Ile Ala Asn Ile Asn Ser Pro Asp Gln
 180 185 190

Ile Val Val Ser Gly Pro Ala Gln Asp Ile Glu Arg Ala Arg Gln Cys
 195 200 205

Phe Val Asp Arg Gly Ala Arg Tyr Val Pro Leu Asn Val Arg Ala Pro
 210 215 220

Phe His Ser Arg Tyr Met Gln Pro Ala Ala Ser Glu Phe Glu Arg Phe
 225 230 235 240

Leu Ser Gln Phe Gln Tyr Ala Pro Leu Arg Cys Val Val Ile Ser Asn
245 250 255

Val Thr Gly Arg Pro Tyr Ala His Asp Asn Val Val Gln Gly Leu Ala
260 265 270

Leu Gln Leu Arg Ser Pro Val Gln Trp Thr Ala Thr Val Arg Tyr Leu
275 280 285

Leu Glu Gln Gly Val Glu Asp Phe Glu Glu Leu Gly Pro Gly Arg Val
290 295 300

Leu Thr Arg Leu Ile Thr Ala Asn Lys Arg Gly Ala Pro Ala Pro Ala
305 310 315 320

Thr Ala Ala Pro Ala Lys Trp Ala Asn Ala
325 330

<210> 16

<211> 417

<212> DNA

<213> Myxococcus xanthus

<400> 16

Met Ser Thr Ser Pro Val Gln Glu Leu Val Val Ser Gly Phe Gly Val
1 5 10 15

Thr Ser Ala Ile Gly Gln Gly Ala Ala Ser Phe Thr Ser Ala Leu Leu
20 25 30

Glu Gly Ala Ala Arg Phe Arg Val Met Glu Arg Pro Gly Arg Gln His
35 40 45

Gln Ala Asn Gly Gln Thr Thr Ala His Leu Gly Ala Glu Ile Ala Ser
50 55 60

Leu Ala Val Pro Glu Gly Val Thr Pro Gln Leu Trp Arg Ser Ala Thr
65 70 75 80

Phe Ser Gly Gln Ala Ala Leu Val Thr Val His Glu Ala Trp Asn Ala
85 90 95

Ala Arg Leu Gln Ala Val Pro Gly His Arg Ile Gly Leu Val Val Gly
100 105 110

Gly Thr Asn Val Gln Gln Arg Asp Leu Val Met Gln Asp Ala Tyr
115 120 125

Arg Glu Arg Val Pro Phe Leu Arg Ala Ala Tyr Gly Ser Thr Phe Met
130 135 140

Asp Thr Asp Leu Val Gly Leu Cys Thr Gln Gln Phe Ala Ile His Gly
145 150 155 160

Met Ser Phe Thr Val Val Gly Gly Ala Ser Ala Ser Gly Leu Leu Ala Val
165 170 175

Ile Gln Ala Ala Glu Ala Val Leu Ser Arg Lys Val Asp Val Cys Ile
180 185 190

Ala Val Gly Ala Leu Met Asp Val Ser Tyr Trp Glu Cys Gln Gly Leu
195 200 205

Arg Ala Met Gly Ala Met Gly Thr Asp Arg Phe Ala Arg Glu Pro Glu
210 215 220

Arg Ala Cys Arg Pro Phe Asp Arg Glu Ser Asp Gly Phe Ile Phe Gly
225 230 235 240

Glu Ala Cys Gly Ala Val Val Val Glu Ser Ala Glu His Ala Arg Arg
245 250 255

Arg Gly Val Thr Pro Arg Gly Ile Leu Ser Gly Trp Ala Met Gln Leu
260 265 270

Asp Ala Ser Arg Gly Pro Leu Ser Ser Ile Glu Arg Glu Ser Gln Val
275 280 285

Ile Gly Ala Ala Leu Arg His Ala Asp Leu Ala Pro Glu Arg Val Asp
290 295 300

Tyr Val Asn Pro His Gly Ser Gly Ser Arg Gln Gly Asp Ala Ile Glu
305 310 315 320

Leu Gly Ala Leu Lys Ala Cys Gly Leu Thr His Ala Arg Val Asn Thr
325 330 335

Thr Lys Ser Ile Thr Gly His Gly Leu Ser Ser Ala Gly Ala Val Gly
340 345 350

Leu Ile Ala Thr Leu Val Gln Leu Glu Gln Gly Arg Leu His Pro Ser
355 360 365

Leu Asn Leu Val Asp Pro Ile Asp Ser Ser Phe Arg Trp Val Gly Ala
370 375 380

Thr Ala Glu Ala Gln Ser Leu Gln Asn Ala Leu Val Leu Ala Tyr Gly
385 390 395 400

Phe Gly Gly Ile Asn Thr Ala Val Ala Val Arg Arg Ser Ala Thr Glu
405 410 415

Ser

<211> 262
<212> DNA
<213> *Myxococcus xanthus*

<400> 17

Met Gln Ala Ala Ser Pro Pro His Arg Asp Tyr Gln Thr Leu Arg Val
1 5 10 15

Arg Phe Glu Ala Gln Thr Cys Phe Leu Gln Leu His Arg Pro Asp Ala
20 25 30

Asp Asn Thr Ile Ser Arg Thr Leu Ile Asp Glu Cys Gln Gln Val Leu
35 40 45

Thr Leu Cys Glu Glu His Ala Thr Thr Val Val Leu Glu Gly Leu Pro
50 55 60

His Val Phe Cys Met Gly Ala Asp Phe Arg Ala Ile His Asp Arg Val
65 70 75 80

Asp Asp Gly Arg Arg Glu Gln Gly Asn Ala Glu Gln Leu Tyr Arg Leu
85 90 95

Trp Leu Gln Leu Ala Thr Gly Pro Tyr Val Thr Val Ala His Val Gln
100 105 110

Gly Lys Ala Asn Ala Gly Gly Leu Gly Phe Val Ser Ala Cys Asp Ile
115 120 125

Val Leu Ala Lys Ala Glu Val Gln Phe Ser Leu Ser Glu Leu Leu Phe
130 135 140

Gly Leu Phe Pro Ala Cys Val Met Pro Phe Leu Ala Arg Arg Ile Gly
145 150 155 160

Ile Gln Arg Ala His Tyr Leu Thr Leu Met Thr Arg Pro Ile Asp Ala
165 170 175

Ala Gln Ala Leu Ser Trp Gly Leu Ala Asp Ala Val Asp Ala Asp Ser
180 185 190

Glu Lys Leu Leu Arg Leu His Leu Arg Arg Leu Arg Cys Leu Ser Lys
195 200 205

Pro Ala Val Thr Gln Tyr Lys Tyr Ala Ser Glu Leu Gly Gly Gln
210 215 220

Leu Leu Ala Ala Met Pro Arg Ala Ile Ser Ala Asn Glu Ala Met Phe
225 230 235 240

Ser Asp Arg Ala Thr Leu Glu Ala Ile His Arg Tyr Val Glu Thr Gly
245 250 255

Arg Leu Pro Trp Glu Ser

B
cont.

<210> 18
<211> 256
<212> DNA
<213> *Myxococcus xanthus*

<400> 18

Met Gly Ile Met Thr Glu Gly Thr Pro Met Ala Pro Val Val Thr Leu
1 5 10 15

His Glu Val Glu Glu Gly Val Ala Gln Ile Thr Leu Val Asp Arg Glu
20 25 30

Asn Lys Asn Met Phe Ser Glu Gln Leu Val Arg Glu Leu Ile Thr Val
35 40 45

Phe Gly Lys Val Asn Gly Asn Glu Arg Tyr Arg Ala Val Val Leu Thr
50 55 60

Gly Tyr Asp Thr Tyr Phe Ala Leu Gly Gly Thr Lys Ala Gly Leu Leu
65 70 75 80

Ser Ile Cys Asp Gly Ile Gly Ser Phe Asn Val Thr Asn Phe Tyr Ser
85 90 95

Leu Ala Leu Glu Cys Asp Ile Pro Val Ile Ser Ala Met Gln Gly His
100 105 110

Gly Val Gly Gly Phe Ala Met Gly Leu Phe Ala Asp Phe Val Val
115 120 125

Leu Ser Arg Glu Ser Val Tyr Thr Thr Asn Phe Met Arg Tyr Gly Phe
130 135 140

Thr Pro Gly Met Gly Ala Thr Tyr Ile Val Pro Lys Arg Leu Gly Tyr
145 150 155 160

Ser Leu Gly His Glu Leu Leu Leu Asn Ala Arg Asn Tyr Arg Gly Ala
165 170 175

Asp Leu Glu Lys Arg Gly Val Pro Phe Pro Val Leu Pro Arg Lys Glu
180 185 190

Val Leu Pro His Ala Tyr Glu Ile Ala Arg Asp Leu Ala Ala Lys Pro
195 200 205

Arg Leu Ser Leu Val Thr Leu Lys Arg His Leu Val Arg Asp Ile Arg
210 215 220

Arg Glu Leu Pro Asp Val Ile Glu Arg Glu Leu Glu Met His Gly Ile
225 230 235 240

B
CMT

Thr Phe His His Asp Asp Val Arg Arg Arg Ile Glu Gln Leu Phe Leu
245 250 255

<210> 19
<211> 424
<212> DNA
<213> Myxococcus xanthus

<400> 19
Met Leu Asn Leu Ile Asn Asn His Ala His Gly Tyr Val Val Thr Pro
1 5 10 15

Val Val Leu Ala Cys Asn Asp Ala Gly Leu Phe Glu Leu Leu Arg Gln
20 25 30

Gly Pro Lys Asp Phe Asp Arg Leu Ala Glu Ala Leu Arg Ala Asn Arg
35 40 45

Gly His Leu Arg Val Ala Met Arg Met Phe Glu Ser Leu Gly Trp Val
50 55 60

Arg Arg Asp Ala Asp Asp Val Tyr Ala Val Thr Ala Ala Ala Ala
65 70 75 80

His Arg Ser Phe Pro Arg Glu Ala Gln Ser Leu Phe Ala Leu Pro Met
85 90 95

Asp Arg Tyr Leu Arg Gly Glu Asp Gly Leu Ser Leu Ala Pro Trp Phe
100 105 110

Glu Arg Ser Arg Ala Ser Trp Asp Thr Asp Asp Thr Leu Val Arg Glu
115 120 125

Leu Leu Asp Gly Ala Ile Ile Thr Pro Leu Met Leu Ala Leu Glu Gln
130 135 140

Arg Gly Gly Leu Lys Glu Ala Arg Arg Leu Ser Asp Leu Trp Ser Gly
145 150 155 160

Gly Asp Gly Arg Asp Thr Cys Val Pro Glu Ala Val Gln His Glu Leu
165 170 175

Ala Gly Phe Phe Ser Ala Gln Lys Trp Thr Arg Glu Asp Ala Val Asp
180 185 190

Ala Glu Leu Thr Pro Lys Gly Ala Phe Ile Phe Glu Arg Ala Leu Leu
195 200 205

Phe Ala Ile Val Gly Ser Tyr Arg Pro Met Leu Ala Ser Met Pro Gln
210 215 220

B1
CONT

Leu Leu Phe Gly Asp Gys Asp Gln Val Phe Gly Arg Asp Glu Ala Gly
225 230 235 240

His Glu Leu His Leu Asp Arg Thr Leu Asn Val Ile Gly Ser Gly His
245 250 255

Gln His Arg Lys Tyr Phe Ala Glu Leu Glu Lys Leu Ile Ile Thr Val
260 265 270

Phe Asp Ala Glu Asn Leu Ser Ala Gln Pro Arg Tyr Ile Ala Asp Met
275 280 285

Gly Cys Gly Asp Gly Thr Leu Leu Lys Arg Val Tyr Glu Thr Val Leu
290 295 300

Arg His Thr Arg Arg Gly Arg Ala Leu Asp Arg Phe Pro Leu Thr Leu
305 310 315 320

Ile Ala Ala Asp Phe Asn Glu Lys Ala Leu Glu Ala Ala Gly Arg Thr
325 330 335

Leu Ala Gly Leu Glu His Val Ala Leu Arg Ala Asp Val Ala Arg Pro
340 345 350

Asp Arg Leu Ile Glu Asp Leu Arg Ala Arg Gly Leu Ala Glu Pro Glu
355 360 365

Asn Thr Leu His Ile Arg Ser Phe Leu Asp His Asp Arg Pro Tyr Gln
370 375 380

Pro Pro Ala Asp Arg Ala Gly Leu His Ala Arg Ile Pro Phe Asp Ser
385 390 395 400

Val Phe Val Gly Lys Ala Gly Gln Glu Val Val Pro Ala Glu Val Phe
405 410 415

His Ser Leu Val Glu His Leu Glu
420

B1
C1
C2

<210> 20

<211> 19053

<212> DNA

<213> Myxococcus xanthus

<400> 20

gtcgacgttg acgtcgcccg gtggcggtgcc gtgtgtcttc ttgcacgcgg aggtgcgcga 60
ggtggcgccggc gacggccggc gcggggccgct gtgtcgcggt gagcgcgcgt atgcgcgggt 120
actggcgctg cgtggccagc gcctccatgc ttccgggtcc tttcgcccg cgtcgtgtat 180
ggctccggtg gaggtgcggcc ggtcaaggg cctgcccaggc acgggtgcggc cgtcgtggta 240
tcagacggcg caccggagg ccctgtccctg ggagcgcgtg ggcgcgggtgg gcaatccctg 300
cctcgtggtg ggtgaactcc ggaggggccc tggcggggc agctacgccc tggcgggtcg 360
ggagggcgcc ccccgatggtggcgttgg accccaggct ccggccacct gtggacgtcg 420

ggcgccggccgg gcctggccggc acttcgcccgc ggccgggggt ctgtccatgg ccggggccgt 480
cgctctgtca gggcgctgt gagacgcgcg gcggggggccg taccggcccg ccagaaacgt 540
gatgcgcgcg caggcctcgc ggtccggcga ctgacgcccgg gcccgtcgg gactcgctca 600
ggcgccctcg gtgcgtcgcg cggggagaa cacgagctgt tccctcgctg ccggccacccg 660
cacgggtgagg gtcccgccca cggccggag gcccagcgcg gtggactcgcg ccagggtccga 720
gagcaggggag cccgcagcgc gcagggggaa gcccgtggc cacatgcctt ccagctcgcc 780
gaacacggcgcg cgcagctgt cggggccctt ggttcgtcgc atggcgcggc gcaggcgcac 840
catgcccgtc acgaagtttc tgccacccat gagcgttgg gggttcttgg agatgaccc 900
cgcaagctg aagtggccgc caccggccgc qactcgctt tcgatgagct gcagggtccctt 960
gccaagggtcg cgcaccgtgg ggtcccttgc gaagtaccac ttggcgtatc gcgcggcgcg 1020
gtctgggtac aagtattca gcatgaggct gccttgc tgcacccgtc ggaggtaggc 1080
ctcccgatg gggcgtaga cccgcgcgcg gaactcgccg gaggcgcggg gaagctgc 1140
gtctccaggcc agcgcgcggg ccagggcgcg tgagaagcgg gacagctcga gctctggat 1200
ggggggcacc agggcgcggg acgagtcatc cgcacatc agctcgagcc gcgtttcgat 1260
gggtgtctt cccatgtctt cctgcatgtt gctgtatggc gtggccgcgt ctttcgtc 1320
cagcacgcgc cagacgcacgg cgtcatccac cgccttgc aggtccgtc gcgcacagctt 1380
cttctcgccgt ccccacacga tcaggtcgcag gccgttagcgc aagtccggg ccacgggggt 1440
gggagagaag cgcgtatgt cccgcgttgcgt gtcacccacc cactggccgc ggggtcccg 1500
cagcttgc tctgtatggcc cccgcgttag gaaacgcgttgcgt ccggggccgc cttggatgtt 1560
cttctcgccgc acccaggact gcttgcgtcgt gtttcgc ttagcgtact cgaacgcac 1620
cagccccagt gagaagcgcg tggccgtctt ggcgttagt gtggggcgt gcaggtatgt 1680
gcccgcgc aactcgctt gctgggggg gaggcttc atccacttca gcagctccac 1740
cagggttgc ttagcggagg actcgtggaa ggcacatcgcg gtgcgttgc ccacgcac 1800
ctccgcgcgc gttggaggctt cccacaggcgc caggtatttc tactcgaagc ctcggcgcac 1860
ctggcgtcgc acggcgctt ccagcgactc tgcacccgcg gcttgcgttgc gggcccgac 1920
ctggcgcgc gttggccggg tccacccgtt cccacccgtt ccaaggcgc ccacggccac 1980
gcccgcgc tttctctgtat tgcgtccatgg cttccgttgc tggcccttgc ccaccaggcc 2040
gtccatgatt tccacccgcg tggccgttgc tttccgttgc tggccgttgc cggggcccg 2100
cagcgccatc agcaacttgc cccatgttgc cttccgttgc gcccggccca gcagctccac 2160
gttgatgc cccgcgcgcg agacgcgcgtt cccatggac ttcacccgttgc gcacggcgcac 2220
ctggatgcgc cccgcacggg gacggggagaa gctgagctgg tagtgcgttgc tgagggacac 2280
ccggccgcac agggccgcct tggccgttgc tttccgttgc cccacccgttgc gcccggccac 2340
gaagccccagg cgggttaggt tggggggaa gatgtccgc cagttgtatggc ccaggccgttgc 2400
gtggagccgc cccgcgcgc acgcacccgc cccgttgcgttgc tggccgttgc ccaggccgttgc 2460
gttgcgttgc atcagccgc acgcacccgc acgcacccgc acggccctac gcacccgttgc 2520
gcccgcgc gtcgcgcgtt agtcgttgc cgttgcgttgc agctccgttgc gtttcgttgc 2580
ggcgaggaaag gacaggctgg cgcacccgc ggcacccgc cccgttgcgttgc tggccgttgc 2640
catccacccgc atgtcacttc ccagcgcacccgc tttccgttgc gggcccttgc cggccgttgc 2700
ccggccgcgc tggccgttgc tggccgttgc tttccgttgc cccacccgttgc gggcccttgc 2760
gatggacgt tggccgttgc tttccgttgc cccacccgttgc gggcccttgc cggccgttgc 2820
ccggccgcgc tttccgttgc tggccgttgc cccacccgttgc gggcccttgc cggccgttgc 2880
ctggcgttgc acgtccggc acgcacccgc cccacccgttgc gggcccttgc cccacccgttgc 2940
gttgaccact ttcacccgttgc tttccgttgc cccacccgttgc gggcccttgc cggccgttgc 3000
tttcgttgc acgcacccgttgc cccacccgttgc gggcccttgc cggccgttgc 3060
cagccgcacccgc atgtcacttc tttccgttgc gggcccttgc cccacccgttgc gggcccttgc 3120
ctggccgttgc tttccgttgc cccacccgttgc gggcccttgc cggccgttgc 3180
ccggccgcgc gggccgttgc tttccgttgc cccacccgttgc gggcccttgc cggccgttgc 3240
gttgacccgttgc cccacccgttgc gggcccttgc cccacccgttgc gggcccttgc 3300
tgacacccttgc acgcacccgttgc gggccgttgc cccacccgttgc gggcccttgc 3360
ggtgccgttgc gggccgttgc cccacccgttgc gggcccttgc cccacccgttgc 3420
gttcccgccac gttcccgccac cccacccgttgc gggcccttgc cccacccgttgc 3480
ggcgccgcgc gggccgttgc cccacccgttgc gggcccttgc cccacccgttgc 3540
gttgacccgttgc tttccgttgc cccacccgttgc gggcccttgc cccacccgttgc 3600
tgccgttgc gggccgttgc cccacccgttgc gggcccttgc cccacccgttgc 3660

gggtggccgc gacgatgcgc acgtccacot tcacggcctg ggtgcctccc acgcgctga 3720
actcgcgatc ctggatgacc cgcagcaact tgccctgcac cggcaggggc agctgccaa 3780
tctcgatcgt gaacacggtg cggccgcgtgg cgcgttcgaa ctgcccggc acgcgggtgg 3840
ccgcgcggta gaaggcgcgg cgttcgtggc ogaagagctc gtttcgtatc agcgtggcgg 3900
gcagcgcgcgc gcaatccacc ttgtatgaaagg gtcgtccct gcggggacca ttacgtgga 3960
cggcacgggc gaacagctc ttgcgcgtgc caacttcgccc ggcagcagc accgtcgatc 4020
cggtggcgcg ggccttcgc accagtccgtt agatggctg gagctgcggg gactgcgcga 4080
tgatgcgggtt gaagaagtag cccacccgtta cctggggctg ctccctcgcg cgctggagct 4140
cttgatagag gtcgtgcgtc tgaggggcgg tgctcaatcg cgaggcgtatc gcggtgagcc 4200
gtcgctgtc ctgcgtggtg aagcggtct cggcggcggc gtggaggacc tggagcacgc 4260
cgttagagggc gccgtccccc tcgcgcgttg gcacggcggag caggctggtg gtgcggtagc 4320
ccgtcatccg gtcgtatcgc gcaagaagc gtcgtcgcc ggcgggttc ggcacgttga 4380
tggcgtccc cgccttcgc acgggtccgg cgcacccctg gcccacgttgc acgcgaatct 4440
gggacaccc gggcagggtc gggcggcgc tgaacacgcgc gggcggggc gggtccagca 4500
gccagagcgt gccgcggcc gcttcgcagg tgatggcgtatc gcggtccatc agcgtctgga 4560
ggaacgcgtc gagggtccacc tcccttcgcg cgcgttcacc gagggggagg aggacccggg 4620
agacgtccga gggggcttgg ggcatggcgg gcaacggcgg caggacaaag gggaggccg 4680
caccataaca tccagaggcc atgggactgc ccccttcag cgcgcggc cccgcaccag 4740
ccgctggcct tcgcgtgcgg gtcgtccacac ggggaagagg ggcggggctt gtcggccat 4800
gtccctcgagc atgtcgatcgc cgtgcgcgg gtcgtgttg aacagcaca gccggcgcgc 4860
ccccaccagc cggccacgc cgcgcgcac catcatggtg gatggccccc agcccttctt 4920
cgccacgcgc ttgcggccct cgtattcgtc cggcgtgtac tgcgtatcga ggcacaggac 4980
gtccgcgcgc tcgaagaggc ggcacccatc cggcgcgagc tccgtccatc gcaatccac 5040
gtccgtggcg tagacgaac aatggccatc cgccttcagg cggtagcaca ggcacccctg 5100
cggtgtccgc acgtcgatgg gcgtagcgc gaaaggggccc acctccatgg gtcggccatg 5160
caacgcggag cggaaatcaca tccggcgaggc catggtgctc agggcaccatc gaaaatgaag 5220
cggtcgatc tgcgcggcca actcggaactg gaggcgttgc gcccatttcg cgcgggacc 5280
gttagagcgtc agctcggtac tggcagccca ggcggcgtg aagaaggggg aggccctgcac 5340
gtggtcccaa tgcagatcg agaagaagag cgtggctcc tggggcgcgc ctgcgtatc 5400
catgatttc cccagtgcgc ggtatggccgt cccgcacatc aggatggcgc ggtggccctg 5460
gtctgttccacc tccacgcagg cgtgttgc accaatgcgc gaggccaca cccggatgtct 5520
cccccaacg ccatgaaacc ggactccatc cgtaaatgttgc cttgtatggg gggccctcgc 5580
ctgggacgcc ctcatgcggc gagecctcaga gacgggggtg tgccattccc aaatgcggg 5640
aatcaggagc gccggcctcg ggctgttccca cgggtgtcc agaatggatc ggcgtccat 5700
ggtcggccgc atccaaacgc gtgcagggtc cccgcaggac gggggggccg gcaatccatc 5760
caacgtccca cggcgtccatc gtcttcagat ctctccatc gggggaaaggc gtccaggagg 5820
ttgcacccgg catcgagcgg ggctgtgttgc ttcagatctt gtcggagctt cggacacaac 5880
cgtctgggtt ctgggaatgc gccggcttcc gttcaatccca gatgttgcata atggcttcgc 5940
agtgcgggtt tagcaatccct cggggcgtaa ccacgcgtt gaaaggcagtc acgcgttcgt 6000
cacgcgttggg gtgtttccag ctcaatcggtt gttatccatc caggggcggtt tgctgtacac 6060
gtccgtccatc ggaagcgatc gcaaaacaat gaaaacgggtg tcgttgcgc ggccttgggc 6120
ctccagaacg ccatccctgc ggcacccaggc agccggaaatt tgagacgggg ctgtcaggccg 6180
tttgcacccca aggtatcgcc ggggggttgc gggcggccccc gaccagaatt cgggtgggtt 6240
gccagttt gtcagattt gagaatagc aggctggggg gaagttgcac tgccctggcc 6300
gcccgtgtgtt gagaacgcattt ggggtgcatt gtcgtccgc gtcacatcacc agaaatggc 6360
tgccgtccatc tggggaaac acggctacgc gtttcgtcc ggcacgtaca cgcctccaa 6420
gtccgtccatc gtcgtccatc cttcccccggg taccccttgc gtcgttacca 6480
gcccgtccatc cggtaccgcgca tcgtccggcgc gcccggggcgttccatc gtcgttacca 6540
cgccggccgcg gaaaggcgtgc cgcacaggatc attggaggccatc ctcggccggg tcggccatc 6600
gggtgtccatc tccatccatc gtcgttgcgtt ggggtgggg caggcgtgc gtcacatcga 6660
agggccctgg acaggctgg aaggaaatgc ggtgcgttgc gaggccaaatc tccgttcat 6720
tgttcgttgc gggctgttgc acgcgtccgtt gtcgttgcgtt gtcgttgcgtt gtcgttgcgtt 6780
accgtatccacc gactgtatcc cgggacatcc cttccatc ttcgttgcgtt gtcgttgcgtt 6840
gcaaggcgttc acggccgtt gtcgttgcgtt gtcgttgcgtt gtcgttgcgtt gtcgttgcgtt 6900

B
Cont

ccagcagtgt tcgggagggtg gtacccgaac tcgaatcaca tccgtcgag ccggaggatg 6960
acctggctgg actggggcgcg aactcgctcg accgcgccga aatcgtaac ctcacgctgg 7020
agaagctggc gctcaacatc cccgggtcg agctgattga cgcgaagacc attggcgggc 7080
tggtggacgt cttcacgcg aggctgttag gegaagccat gggccggc gggattgaag 7140
ccatgaatgc tctatgtggc atcgccaggt tggatgtgtt gcagctggcg acccaccgtg 7200
gcctggacac ctccgccttc gcgaaacctgc tcatggagga gaagaccgtc ccgcctccct 7260
atgaggaccc tgcacccatc ggcgtgaatg ccgcctggcc catctggac cagtggaccg 7320
cggcggaaacg ggacagcatc gagctgtgg tggatgtgcac ggagtccctcg ttgcacttcg 7380
gcaaggccat gaggcaccatc ctgcaccaggc acctggggct gaggccgaac tgccggctca 7440
tcgagctcaa gagcgccttc tactccgggg tcgcggggct gcagatggcc gtcaacttca 7500
tcctgtcggc cgtgtcggcc ggggccaagg ccctgggtt ggcctccgac ctgtcgccgt 7560
tctccatcgc cgaaggggga gatgcctcca cggaggadgt gtccttcggc gaggccgagct 7620
cggtgtcggg cgccgtggcc atgctgtga ggcacacgccc cgggtgttc cgcgtcgacg 7680
tggggggcggaa cggctactac ggctacgagg ttagtggatad ctgcgcggcc gtggcggaca 7740
gcaagcggg agacgcggac ctgtcgctcc tctcgatctt ggactgtgtt gagaacgcct 7800
tccgggagta caccgcgcgc gtcccccggc cgaactacgc ggagagcttc ggatccctcg 7860
cctccacac gccgttggc ggcatggta agggcgccca ccgcacatg atgcgaagt 7920
tctccggcaa gaaccgcggg gacatcgaag cggacttcca gcggcggatg gccccgggc 7980
tgacctactg ccagcgcgtg gggaaacatca tggcgcgac gatggcgtc tcgccttcg 8040
ggaccatcgaa ccacggcgc ttgcaccccg cgaaggatg tggatgttc tcgtatggct 8100
cggtgtcag ctcggagttc tttagccggc tggatgtggc ggaaggccag cagccgcgc 8160
gcgccttggc gctgggagaa ggcgtggggc gccggcagca gtcgcctatc cggattacg 8220
acgcgcgtc gaaaggaaac ggcctgggtc gcttggggac cggaaacgc gactgtgatt 8280
tcggatgtgtt cggcagcatc cggccggggc ggtggggcag gcccgttgc tcctgtcgg 8340
cgattcgtga ctccatcgc gactaccaat ggatcccttgc gcttcggggc ttcgagcaaa 8400
gcatatcgca gcttagcgcac ggccgtcccc ctgacggccc gtgacagegc ggtgagccgc 8460
cggtcgcaaa tcaccccaatc catgtcgccg cagacgtctt tggatgtggc gcaatggc 8520
gactggcat gggacaccgt cagccgcctt tggatgtggc acgtgttgc cgcgaccaac 8580
gcctcaggcg cgcgccttgc cttggccatc cttggccatc tattacttc gcatccgggg cgcgcgcgc 8640
ctgcaccccg ggcgcgtcg cttggccatc acgtggacg tcaatgttgc ggtgtacaac 8700
ttccggcagcc aatccgttgc gacggatgc cgcacatgc acagccggaa gggggggcgt 8760
ccggaggccg atgccttcgg ccatgaagag ctgtacgacg agcccgagcc agggccgcac 8820
tacgcggaga cttcaacccg tggatgtacgc gcttcggacg gcaatgttgc gcaatgttgc 8880
atcaatgttgc cgcgcgtgg gttccatgtac gcacaccgtc cgcgttgc ggcacatgc 8940
tcgcgcgc gggcctatgg ggacgcgcgg ggcggggca ctttcacgc tggatgttgc 9000
gcggagttacc ggctgaccgt ggacccgttcc cgcgttgcgt acgcgggttgc cgtcatccgg 9060
gacgtcaatgg gggggggctt catctacttc ggtgttgcatt ttcgtatggt ggactggcc 9120
atctggcagc tggcgaggca ccaggacgc acgcggcagg ctttcgttgc ggcgtgggt 9180
ctggaccagc aactgtgtt cctcgccatc gccggcgttgc acaccatcc cgcacatgc 9240
gtgcagcaatc gggagccggg gggccgggg gaagagctgt tcaacgttgc gatgcgcgc 9300
ggccgcgcagg ggcgggacat cggcggtggc acgttgcagg tggatgttgc cgcgcgttgc 9360
gaaggaggcc ggcgtgggtt ggcgttgc acgcggatgc agacgaacaa atccggggag tggatgttgc 9420
gtccatcgatc cgcgttgc cccgcgttgc ctccacgc acgtggggcc actgttgc 9480
ccgcggatgttgc ggcgcggact cgcgttgc acgtggggcc ttcacgttca tccgttgc 9540
ccttcggccatc cagaagacgc cactggcgaa tggatgttgc acaccatcc cgcacatgc 9600
ggatgttgcgttgc tggccgggtt ggttgcgttgc acaccatcc cgcacatgc 9660
tcgaggccatc caacgcctac ggcggccccc cttccatcc ggttgcgttgc tggatgttgc 9720
gcccggcgttgc ggcggccatc aacgttgc acgtggggcc ttcacgttca tccgttgc 9780
tcgcgcgttgc aggacccgttgc cacaacgc acgttgc acaccatcc cgcacatgc 9840
atgtcgccatc aggccggaa ggcgttgc acaccatcc cgcacatgc 9900
gacttcaggca agtccatcc ctcgtatgttgc acgtggggcc ttcacgttca tccgttgc 9960
cggttcctgg aggttgc acgttgc acaccatcc cgcacatgc 10020
gcgttgc acgttgc acaccatcc cgcacatgc 10080
acgcgttgc acgttgc acaccatcc cgcacatgc 10140

B
Cont

gtgtctgtgg ggcgacgagcc ggcgtgtatcgg aagatggacc tgggagcggtt cggcaactac 10200
agctacgacg tcttcgacac cgcggggccc tcgcccggaga ttgatatcgg cgacgtggac 10260
cggtcgctct tcacgtacct ggactgcgtc aagcacagct tcgcccgta tggccggccg 10320
gtggacgttg tcgactctgt gtcgacgttc gactaccctgg cgatgcacac gccgttcg 10380
ggactgttga aggccgggca cgcgaagatg atgcgcgagc tcacccctgt cgacgtggac 10440
gaaatcgaag cggacttcgg cggccgtgt aagccgtac tgcagtaccc gagtctggc 10500
ggaaacctgt gctccggcgtc cgtgtacccgtl agccgtgtca gcatcatcga caccatcaag 10560
cccgagcgtt cgcgtcggtt gggatgttc tcctatgggt cgggttgcgc tcggaggatc 10620
ttcagcggcg tcacggccc ggagtccgt tcgcgttag ctgggttggc catcggtggc 10680
cacctccggg ggcgcggcca gtcacgttc gaccaatatc tgcattgtc gaaagagaac 10740
cttcgtgtc tggttcaac gaagaaccgg gacgtggacg tggaggccta cttcccg 10800
gtgacgccc gggcggccg cccgcgtatc tcgcgttc gagggtgtl ggactatcat 10860
cgtcagtagc agtgggtgt tgcatacgc caccgttcaat tccgacaaat gaaacacttc 10920
tccgtacga actggccgtc cgcgtgggc tatctcttg cgttgggg cgcatgttc 10980
gcccgcgtc aagtccacaa acagatggc cgcgcacgggg cgaaaaggcc cgtcgccgtc 11040
ttcgatagct ggtggactt ccactacgtg gagaaccgg cgggtcggtt cggctgttc 11100
tccagttcg gcaagaggt ggcgtatcctt tcctctatc tgcgtggccg catctgcac 11160
gtgttgcgtg tggactacta ctctacacg cgcgcgtac gtaagcttca cgcgtggc 11220
ctggcgacgtc tgattggccg cgcgtgggc aactacgtgg accgggtggt cctgcgtac 11280
gtggtgatt tgcgtatc gacgtgggg gaccgttctt attggccctc cttcaacatc 11340
gcccgcacag cggtagtgcgtt agggccgccc ctgtatgttcc tggagtcgtt cgcgcggcc 11400
cgtcagcgt tgcgtcccg ataggccccg ccatgggtgt cgggtccgcg cccggccaa 11460
ggactggagt tcatggggac ctcagagcca gttgaggccg accacccctt gtcaaaacca 11520
ccgcgtgtc cgcgttgcg cgcgttgcg cgcgttgcg gtcggcaat gcccggcattc 11580
gcccgcgttca tgcgttgcgtt ctcggccccc acggagttcc tggacccgtt cgcgcggcc 11640
tacggtgaca ctttcacccat caagattccg gggacgcgc cgttcatcca gaccagcgat 11700
ccgcgttgcgttca ctcgggtatc tttcaagggtt gacccggacc ttttcgttccggg aggaaaggcc 11760
aacaacgggt tgaagccgtt ggtgggttag aactcgttc tggatgttgc cgggaaaggcc 11820
caccggcggt atcgcacgtt catcatgccc accttcgttgc tgcacggat gcatgttat 11880
ggctcggtca tccgggacat cgtcaatcgc ggcgttgcacc ggtggccgtt cgggaaaggcc 11940
ttcgtcggtcc atgaagagac gacgcacatc atgcgtgggg tgattcttcg ggtgattttc 12000
ggccgtggagg acgcccggac cattggccatc ttccggcacc acgtgcacca ggtgtcaag 12060
ctggccctgt tccgttccg gaacggggag ggcaaggccg cgcgtgggg cttcgcgttgc 12120
gcccgtggca aggcgtttcc cttccgttgc gtttgcgttgc cgttgcgttgc gtttgcgttgc 12180
atcatctacc aggagattca ggacccggccg agccaggacatc tgcgtggccg gtcaggacgt 12240
cttcgtgtca tgcgttgcgttgc gcaactacgc acgggttccg tgcgttgcgttgc ctcggagatc 12300
cgcgttgcgttgc tgcgttgcgttgc gtttgcgttgc gtttgcgttgc gtttgcgttgc 12360
tggatgttgcgttcc accacccatc ctcgggttgc gtttgcgttgc gtttgcgttgc 12420
ggccgttgcgttcc accacccatc ctcgggttgc gtttgcgttgc gtttgcgttgc 12480
gatgttgcgttgc tcaaggagac gatgttgcgttgc acggccgttcc tgcgttgcgttgc gtttgcgttgc 12540
ctcaaggagac ctcgggttgc gtttgcgttgc gtttgcgttgc gtttgcgttgc 12600
aacaatctacc gtcaggacatc tgcgtggccg cttccgttgc gtttgcgttgc 12660
gagcgttcc tggaggagac ggttgcgttgc tttccactact tcccttcgttgc gtttgcgttgc 12720
cggaaatgttca tccgggttgc gtttgcgttgc gtttgcgttgc gtttgcgttgc 12780
gttgcgttgc tgcgttgcgttgc gtttgcgttgc gtttgcgttgc gtttgcgttgc 12840
aacaatctacc gtcaggacatc tgcgtggccg cttccgttgc gtttgcgttgc 12900
accgttgcgttgc cccaccatc ctcgggttgc gtttgcgttgc gtttgcgttgc 12960
ggccgttgcgttgc gtttgcgttgc gtttgcgttgc gtttgcgttgc 13020
ggagctgttgc tttccactact tcccttcgttgc gtttgcgttgc gtttgcgttgc 13080
caaggcgttgc tttccactact tcccttcgttgc gtttgcgttgc gtttgcgttgc 13140
ggatccgttgc gtttgcgttgc tttccactact tcccttcgttgc gtttgcgttgc 13200
ggccgttgcgttgc cccaccatc ctcgggttgc gtttgcgttgc gtttgcgttgc 13260
cgagggttgcgttgc gtttgcgttgc tttccactact tcccttcgttgc gtttgcgttgc 13320
ggccgttgcgttgc gtttgcgttgc gtttgcgttgc gtttgcgttgc 13380

tcacgaactgaa gggccgtcc gggatggcg aggtgcagc 13440
catcaactac ccgtcaact tcgtccgtc ggccggatgag gccccctgg gacggattca 13500
gcaggaactc tcccaacgct cggggcggtt ccaccgggtt cccgtcgct acccccttca 13560
ttccctcgac ctggaccccg tgagggagga gtacccgaagc cccgtcccg cgattcgct 13620
gacgtggccg cgaatccccca tgactctgt caccaccggc aaccgggtgc acgacctcg 13680
cagcgaccac ttctggaaacg tggcccgccgccccatccag ctgtacgaca cccgtcgac 13740
actggaggggg cagggcggtt cgcacttcat cccggccgtt cccgtcgac 13800
catcatcaag cgcacccctcg cccggggactc cacgtacagg ctctccctg tgctcagccc 13860
ttctcccgca tcgacccggg gtcgatggg gtacccggg gtcgcgcgt gacggaggcg 13920
cccgacccca gggcccgctc gcaggtggccg cccggcccg gtcgcgcctg ggcgcgtc 13980
acccgaggag cggcgacccg gccgggtaaat gcccggcaagg cccgcgtctt cccggggcg 14040
ggctcgagg agcggccat gggggcccg ctgtacgact agtcccgga cctgacggac 14100
atcgccgaccc ccatctggg gtattccatc aacgtctt gttggagga cccaggcaag 14160
gagctggcgc agacgcagtt caccggccg gctgttacg tggtaacgc gtcagactac 14220
ctgaaggccg tgcgtgaagg agcggaggcag cccgcgttcg tcgcggccca cccgtggc 14280
gagtaacacg cgcgtgtt cccggggcc tcgcacttgc agacgggact gggctggt 14340
aagccggccgg gcgaaactcat gacccggccg tccggaggg cccaggcaag 14400
tgtgtatggcc tggccgttga acaggctt cccggccgtc agctgaccag tctggatatc 14460
gccaacatca actcgcccgca ccagattgt gtcgcgttgc cccggccgggcatc 14520
gcacggcagt gttctgttgc cccggccgtc cccgttgc acgttgcgttgc 14580
tttactcgcc gtcacatca gccggccggc acggacttgc agtgcgttgc 14640
cagtaacgcgc cccgttgc cccgttgc acgttgcgttgc 14700
gacaacgtgg tgcaggggctt ggcctgttgc cccggccgggcatc 14760
gtccgcgtacc tccatccatca gggccgttgc gacttgcagg agtgcgttgc 14820
ctgaccccgcc tcacccatccatca gaccaagccg ggcggccccc cccggccggcc 14880
gcaagttggg cgaatgttgc acggccgttgc cccggccgggcatc 14940
gtccgcgtcc tgagaccactt gatgtccatc ctcacccgttgc cccggccggcc 15000
gttcggggcc tccatccatca tggccaggg ggcggccgttgc acgttgcgttgc 15060
gggccggccca cgggtccggg tggatggagcg gccggccgttgc cccggccggcc 15120
gacgacggcc caccctgggg cggaaatcgc ctcgcgttgc gtcggccggcc 15180
acaactgtgg cgcgtccatca cccgttgc cccggccggcc 15240
ctggaacccggc cccggccgttgc cccggccggcc 15300
caccacgttgc cccggccgttgc cccggccggcc 15360
tttctgtgg cccggccgttgc cccggccggcc 15420
gcacgttgc cccggccgttgc cccggccggcc 15480
gtccggccgttgc atccaggccg cccggccgttgc cccggccggcc 15540
cgtggggccgttgc cccggccgttgc cccggccggcc 15600
gatggccacc cccggccgttgc cccggccggcc 15660
gatggatggcc tccatccatca gggccgttgc cccggccggcc 15720
cgcgtccatca gggccgttgc cccggccggcc 15780
cgcgtccatca gggccgttgc cccggccggcc 15840
gcccacccggc cccggccgttgc cccggccggcc 15900
ggtcaacacc acgaagtcac tccatccatca gggccgttgc cccggccggcc 15960
catcgccac cccggccgttgc cccggccggcc 16020
catcgccac cccggccgttgc cccggccggcc 16080
cccgattgtatccatccatca gggccgttgc cccggccggcc 16140
cgcgtccatca gggccgttgc cccggccggcc 16200
cgcgtccatca gggccgttgc cccggccggcc 16260
cgcgtccatca gggccgttgc cccggccggcc 16320
acaacacccat cccggccgttgc cccggccggcc 16380
agcgtccatca cccggccgttgc cccggccggcc 16440
ttcggccat cccggccgttgc cccggccggcc 16500
gttacccggccat cccggccgttgc cccggccggcc 16560
gcaaggccaa cccggccgttgc cccggccggcc 16620

cgagggtcca gttcagtc tccgagctgc tttcgggt gttcccgcc tgcgtatgc 16680
cgttcctcgc cggcgaaatc ggcattcaggc gggcgcaacta cctgacgctg atgacgcggc 16740
ccatcgacgc gggccaggcg ctgagctggg ggttggcgga cgccgtggac gccgatagcg 16800
agaagctttt gcccgtccac ttgcgcaggc tgcgggtcct gtcgaagcca gccgtgaccc 16860
agtacaagaa gtacgcctcc gagctggcg gccagctgct cgcggccatg ccccggtcca 16920
tctccgcacaa tgaggcgatg ttctccgcac ggcacacgct ggaagccatc catcgctacg 16980
tggagacagg ccgactccca tggaaatcat gacggaagga acgcaatgg cggcggtgg 17040
cacgcctcat gaggtggagg aggggtggc gcagatcacc ctgggtggatc gcgagaacaa 17100
gaacatgttc agcgagcgc tcgtcgccaa gtcatcacc gtgttgcgcgaa aggtgaatgg 17160
aaacgagcgc taccgcggc tggtgtcgc cggctacgac acctactcg cgctcgccgg 17220
gaccaaggcc ggcctgtgtt ccatctcgca cggcatcgcc tccttcacac tcaccaactt 17280
ctacagcctc ggcgtggagt ggcacatccc ggtgatttcc gcatcgagg gacatggcgt 17340
aggcgccggg ttgcgtatgg ggctgtcgccggactcgatg gtcttgcgcg 17400
ctacacgacg aacttcatgc gctacggctt cggccgggg atggcgcca cgtacatcg 17460
gccgaagcgg ctgggtact cgctcggca tgagctcctg ctcaacgcca ggaactaccg 17520
cggcgccgac ctggagaagc gggcggtgc ttccgggtt tgccgcgcgaa aggaagtcgt 17580
gccccacgccc tacgagatgg cgagggaccc ggtcgcgaaa cctcggtgt cgctcggtac 17640
gctcaagcgg cacctggcgc ggcacatccg cggagacgtt cggacgtca tcgagcgtga 17700
gctggagatg cacggcatca cttccatca cgacgacgtg aggaggcgca tcgagcagct 17760
gttcctgtt ggcgcgcggc tatgttaac ctgatcaaca accacgcaca cggttatgt 17820
gtcacgccc tggccctggc ctgcaacgac gtcggctgt tcgaaactctt gggcaggga 17880
ccgaaggact tcgaccgggtt ggccggaggca ttgcgtgcac accggggaca tctgcgcgtc 17940
gcgtatggatg ttgcgtatgc gctcggtgg gttcgcgcg acgcccgtatg cgtgtacgcg 18000
gtgacggccgg cggcgccgcg gcatcggtcc ttcccgccg aggccgcgtc gctctcgcc 18060
ctgcccatttgg accggatctt ggcggggagg gacggccgtt ccctggcgcc gtgggtcgag 18120
cgctctcggtt cgtcggtggg taccgtatgc acgctgggtgtc gcgagctgtc cgacggcgcc 18180
atcatcacgc cgctgtatgcg cgcgtgggg cctcaaggaa ggcggccgt 18240
ctgtccgacc tgggtccgg gggggatggaa agggacacgtt gctcccgca gggcggtccaa 18300
cacgagctgg cgggttctt ctccgcgcg aagtggacgc tgaggacgc gtcgcacgcg 18360
gagctcacgc ccaaggccgc ctcatctt gageggcat tgctttcgatc catcgccggc 18420
tcgtaccggc cgatgtggc cagcatcgcc cagctgttctt tgggtactcg cggccagggtc 18480
ttccggccgg acgaaggccgg ccacgaacttggacc gaaacccttca cgtgtatggg 18540
agccggccacc agcaccggaa gtacttcgcg gagctggaga agctcatcat caccgttcc 18600
gatgtccgaga acctgtccgc acagccgcg tacatcgccg acatgggttgcggtacggc 18660
acgctccctgtt acgggggttta tgaaacgggtt ctccggcaca cgcggccgggg aaggccgtc 18720
gaccgggttcc gctcacgtt catcgccgcg gacttcaacg agaaggccgt cgaagccgt 18780
ggccggccgc tggccgggtt ggagcacgtt gcttgcgcg cggacgtggc gccggccggac 18840
cgctctatcg aggacctgcg ggcgcgcggg ctggccgcgcg ctgagaataac gtcgcacatc 18900
cgctcggttcc tggaccacgca cctgccttac cagccctcccg cggacaggccg ggggttccac 18960
gccccggatcc ggtcgatcc ggtgttgcgtt ggcaaggccgg gccaggaggtt ggttccggccg 19020
gagggttcc acggcctggt ggagcacctc gag 19053